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Result
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Maximum Match 100%
Listing first 45 summaries
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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3: /cgn2_6/ptodata/1/laa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/laa/6B_COMB.pep:*
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2 US-08-356-786-4

2 US-08-356-786-10

US-08-273-146-47

US-09-648-067A-12

US-09-648-067A-12

US-08-875-811-41

US-08-875-811-49

US-08-875-811-49

US-08-875-811-64

US-08-273-146-53

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US-08-290-592E-20	US-09-943-906-72	US-09-550-374-72	US-09-036-579-72	US-08-713-939A-72	US-09-560-761B-20	US-09-134-000C-5010	US-09-902-540-14530	US-08-964-690-26	US-08-202-047-26	US-09-328-352-5739	US-09-270-767-42252	US-09-248-796A-22117	US-08-484-537-51	US-08-484-537-50	US-08-487-200-51	US-08-487-200-50	US-08-474-040-51
Sequence 20, Appl	Sequence 72, Appl	Sequence 72, Appl	Sequence 72, Appl	Sequence 72, Appl	Sequence 20, Appl	Sequence 5010, Ap	Seguence 14530, A	Sequence 26, Appl	Sequence 26, Appl	Sequence 5739, Ap	Sequence 42252, A	Sequence 22117, A	Sequence 51, Appl	•	Sequence 51, Appl	Sequence 50, Appl	Sequence 51, Appl

RESULT 1

ALIGNMENTS

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; Sequence 26, Application US/08483749A ; Patent No. 6054561 ; GENERAL INFORMATION:
                                                                                                                        US-08-483-749A-26
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                                                           Query Match
Best Local
                                          Matches
                                                                                                                                                                                                                                                       FILING DATE: 07-UN-1995

CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: SAVEREIDE, PAUL B.
REGISTRATION NUMBER: 36,914
REFERENCE/DOCKET NUMBER: 0508
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 601-2585
TELEPHONE: (510) 655-3542
THEODERICATION: TOTALEPAX: (510) 655-3542
                                                                                                                                                                                                                       TELEFAX: (510) 655-3542
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94662-8097
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE SYSTEM: PC-DOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/483,749A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: CHIRON CORPORATION
STREET: INTELLECTUAL PROPERTY -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: ANTIGEN-BINDING SITES OF ANTIBODY TITLE OF INVENTION: MOLECULES SPECIFIC FOR CANCER ANTIGENS
                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: RING, DAVID B.
                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                    LENGTH:
1 LOYALFPYT 9
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Similarity 100.0%; E
9; Conservative 0;
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                                          Score 49; DB 3
Pred. No. 0.16;
Mismatches
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RESULT

US-08-133-804-6

GENERAL INFORMATION:

Huston, James S.

6, Application US/08133804 o. 5534254

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RESULT 3
US-08-461-838-6
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; TOPOLOGY: linear
, MOLECULE TYPE: protein
US-08-133-804-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.25
SOFTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,804
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application US/08461838 Patent No. 5753204
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                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 617-248-7100 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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LENGTH: 243 amino acid
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TITLE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                              APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynth
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
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Local Similarity 100.0%; Pred. No.
les 9; Conservative 0; Mismatch
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ZIP: 02109
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                                                                              COUNTRY:
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Exchange Place, 53 State Street
                                                                              USA
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Exchange Place, 53 State Street
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Houston, L. L.
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                                                                                                                                                                                                                                    Biosynthetic Binding Proteins For
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Best Local Similarity
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Patent No.
                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,386
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthe
TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 243 amino acid
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                                                                                                    INFORMATION FOR SEQ ID NO:
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MOLECULE TYPE:
}-461-386-6
                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Boston
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                                 TOPOLOGY:
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                                               amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Testa, Hurwitz & Thibeault/Patent Department Exchange Place, 53 State Street
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                                                                                                                    617-248-7100
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                                                                                                                                      617-248-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Biosynthetic Binding Proteins
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Pred. No. 0.34;
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Query Match

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Patient No. 5877305
GENERAL INFORMATION:
APPLICANT: Huston,
APPLICANT: Oppermar
APPLICANT: Houston,
                                                                                                                           US-08-356-786-10
                                                                                                                                                                                                                                                                                                                                         US-08-356-786-4
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                                                                                     Sequence 10, Application US/08356786 Patent No. 5877305
                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08356786 Patent No. 5877305
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 07/831,967
PILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
REGISTRATION NUMBER: 27,829
REFRENCE/DOCKET NUMBER: CRP-053
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATING-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OP INVENTION: Biosynthetic Binding Protein for Cancer
TITLE OP INVENTION: Marker
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                 / Match 100.0%;
Local Similarity 100.0%;
nes 9; Conservative 0
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les 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                              222
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                                                                                                                                                                                              LOYAIFPYT 230
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                                                                                                                                                                                                                                                                                                                                                                                                           243 amino acids
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nilarity 100.0%;
Conservative 0
Oppermann, Hermann
Houston, L. L.
Ring, David B.
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                                                     Huston, James S.
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                                                                                                                                                                                                                                                                                 Score 49; DB 2
Pred. No. 0.34;
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                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                    DB 2; Length 243;
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RESULT 7
US-08-273-146-47
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/831,967
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CRP-/
TELECOMMUNICATION INFORMATION:
TELEPAX: (617) 248-7100
TELEPAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 5855885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 47,
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APPLICANT:
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APPLICANT:
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OURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/356 77 FILING DATE:

CLASSIET CLASSIET PC-DOS/MS-DOS

PC-D
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kenten, John H.
APPLICANT: Mark T.
APPLICANT: Titmes, Richard C.
APPLICANT: Williams, Richard O.
APPLICANT: Williams, Richard O.
TITLE OF INVENTION: The Isolation and Production of
TITLE OF INVENTION: Catalytic Antibodies using Phage Technology
NUMBER OF SEQUENCES: 71
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ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault
STREET: Exchange Place, 53 State Street
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                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                            COUNTRY: U
ZIP: 20852
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CANT: Smith, Rodger
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)GY: linear
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                                                                                                                                                USA
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Chiswell, David
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fitzgerald,
                                                                                                                                                                                                                                                                                     IGEN, Inc
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Sequence 12, Application US/09648067A

Patent No. 6627196
GENERAL INFORMATION:
APPLICANT: Baughman, Sharon A.
APPLICANT: Shak Steven
TITLE OF INVENTION: Dosages for Treatment wi
FILE REFERENCE: P1775R1
CURRENT APPLICATION NUMBER: US/09/648,067A
CURRENT FILING DATE: 2000-08-25
CURRENT FILING DATE: US 60/151,018
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Best Local Similarity
7; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REPERENCE: P1775R1
CURRENT APPLICATION NUMBER: US/09/648,067A
CURRENT FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,018
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/213,822
PRIOR FILING DATE: 2000-06-23
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TELEFAX: 301-230-0158
INFORMATION FOR SEQ ID NO: 47:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Baughman, Sl
APPLICANT: Shak Steven
                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 107
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Ryan, John W.
REGISTRATION NUMBER: 33,771
REFERENCE/DOCKET NUMBER: 099
TELECOMMUNICATION INFORMATION:
TELEPHONE: 301-844-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 14-JU CLASSIFICATION: 43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81
                                                                                                                                                                                                                               90 QYYIYPYT 97
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GY: linear
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                                                                                                                                                                                                                                                                  QYAIFPYT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dosages for Treatment with Anti-ErbB2 Antibodies
                                                                   Dosages for Treatment with Anti-ErbB2 Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sharon A.
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77.8%;
                                                                                                                                                                                                                                                                                                                     73.5%;
75.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                    1; Mismatches
                                                                                                                                                                                                                                                                                                                    Score 36; DB 4;
Pred. No. 28;
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25;
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LOYDEFPYT 97

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PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-06-23
NUMBER OF SEQ ID NOS: 15
SEQ ID NO 12
LENGTH: 107
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US-08-737-129A-8
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                                                                                    ; MOLECULE TYPE: US-08-737-129A-8
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GENERAL INFORMATION:
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                             Matches
                                                       Query Match
                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 215 amino acid
                                                                                                                                                                                                                            FILING DATE:
ATTOMEY AGENT INFORMATION:
ANAME: WATTON Cheek, Jr.
REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ikuo FUJ
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
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                          Local Similarity res 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM:
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                                                                                                                              STRANDEDNESS:
                                                                                                                                             LENGTH: 215 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: No. 5885816ember
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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6; Conservat
 1 LOYAIFPYT 9
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805 Fifteenth
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                                                                                                               ESS: single
linear
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JENTION: CATALYTIC ANTIBODIES ENANTIOSELECTIVELY
JENTION: HYDROLYSING AMINO ACID ESTER DERIVATIVES
                                Conservative
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75.0%;
                                            73.5%;
77.8%;
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Street, N.W., #700
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                                Score 36; DB
Pred. No. 54;
0; Mismatches
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US-08-875-811-49
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                                                                                                                                                                                                Sequence 49, Application US/08875811 Patent No. 6045793
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Best Local 9
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Patent No. 6045793
                                                              APPLICANT: Rybak, Susanna M.
APPLICANT: Newton, Dianne L.
APPLICANT: Boque, Lluis
APPLICANT: Boque, Lluis
APPLICANT: Wlodawer, Alexander
TITLE OF INVENTION: Recombinant Ribonuclease Proteins
NUMBER OF SEQUENCES: 64
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APPLICANT: Rybak, Susanna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
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APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,811 FILING DATE: 19-FEB-1998
CLASSIFICATION: 435
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APPLICANT: Boque, Lluis
APPLICANT: Wlodawer, Alexander
TITLE OF INVENTION: Recombinant Ribonuclease Proteins
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Faria, Sugan K.
REGISTRATION NUMBER: 41,7
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ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
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MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 94111-3834
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Similarity 77.8%;
7; Conservative
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Dianne L.
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Pred. No.
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APPLICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Faris, Susan K.
REGISTRATION NUMBER: 41,739
REFERENCE DOCKET NUMBER: 015280-24410(
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1876-0300
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US-08-875-811-64
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APPLICANT: Rybak,
APPLICANT: Newton,
APPLICANT: Boque,
APPLICANT: Wlodawe
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
APPLICATION NUMBER: US/08/875,811
FILING DATE: 19-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                         CITY: San Francisco
STATE: California
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Similarity 77.8%;
7; Conservative
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VENTION: Recombinant Ribonuclease Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                          Newton, Diani
Boque, Lluis
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Pred. No. 87;
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                                                                                                                                       Version #1.30
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/88/875,811
FILING DATE: 19-FEB-1998
CLASSIFICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Faris, Susan K.
REGISTRATION NUMBER: 41.739
REFERENCE/DOCKET NUMBER: 015280-244100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
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Patent No. 6045793
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INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200
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APPLICATION NUMBER:
FILING DATE: 21-FEB-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Boque, Lluis
APPLICANT: Wlodawer, Alexander
TITLE OF INVENTION: Recombinant Ribonuclease Proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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REGISTRATION NUMBER: 41,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Protein
LOCATION: 1..355
OTHER INFORMATION:
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les 7; Conservative
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Boque, Lluis
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77.8%;
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Pred. No. 87;
0; Mismatches
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Best Local :
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                                                                                           Query Match
                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: FAT18, SUSAN K.
REGISTRATION NUMBER: 41,739
REFERENCE/DOCKET NUMBER: 015280-244100US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                     TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Newton, Dianne APPLICANT: Boque, Lluis APPLICANT: Wlodawer, Alexantrus: Recording OF INVENTION: Recording OF SEQUENCES: 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 358 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                          MOLECULE TYPE: protein
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TYPE: amino acid
TOPOLOGY: linear
                                                            Local Similarity nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 19-
CLASSIFICATION:
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                                                                                                                                                                                         ENGTH:
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207 LOYDEFPYT 215
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Similarity 77.8%;
7; Conservative
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                                                                                                                                                           : 358 amino acids
amino acid
GY: linear
                       LQYAIFPYT 9
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                                                             Conservative
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                                                                              73.5%;
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 Mismatches

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                                                                                Score 36;
Pred. No.
                                                                  Mismatches
                                                                                   DB 3;
87;
                                                                                                Length 358;
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Search completed: April 4, 2005, 16:00:52 Job time : 4.60064 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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146.032 Million cell updates/sec
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1: /cgn2_6/ptodata/1/1aa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/1aa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/1aa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/1aa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/1aa/backfIles1.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-08-273-146-45
US-08-273-146-53
US-08-483-749A-26
US-09-726-219A-267
US-08-466-886-27
US-08-466-886-27
US-08-469-617-27
US-09-943-906-74
US-09-943-906-74
US-09-943-906-74
US-09-943-906-77
US-09-627-218B-1
US-08-461-838-6
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US-08-356-786-10
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US-09-195-391-3
US-09-195-391-3
US-08-469-617-26
US-08-469-617-26
US-08-469-617-26
US-08-469-617-26
US-09-144-796A-2076
US-09-144-96A-2076
US-09-144-96A-2076
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                          Sequence 45, Appl
Sequence 53, Appl
Sequence 26, Appl
Sequence 27, Appl
Sequence 74, Appl
Sequence 6, Appli
Sequence 6, Appli
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Sequence 10, Appli
Sequence 3, Appli
Sequence 10, Appli
Sequence 26, Appli
Sequence 27, Appli
Sequence 10, Appli
Sequence 26, Appli
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Sequence 10, Appli
Sequence 27, Appli
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        Sequence
                                                                                       ; TYPE: amino a
; TOPOLOGY: lir
; MOLECULE TYPE:
US-08-273-146-45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1
US-08-273-146-45
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GENERAL INFORMATION:
Query Match
Best Local Similarity
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APPLICANT:
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ALIGNMENTS

Rodger

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APPLICANT: McCafferty, John
APPLICANT: Chiswell, David
APPLICANT: Darsley, Michael J.
APPLICANT: Fitzgerald, Kevin
APPLICANT: Fitzgerald, Kevin
APPLICANT: Wartin, Mark T.
APPLICANT: Martin, Mark T.
APPLICANT: Williams, Richard C.
APPLICANT: Williams, Richard O.
TITLE OF INVENTION: The Isolation and Production of
TITLE OF INVENTION: Catalytic Antibodies using Phage Technology
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS: 105.
                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Vercurrent Application DATA:

APPLICATION NUMBER: US/08/273,146
FILLIG DATE: 1.4-UUL-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: RYAN, John W.
REGISTRATION NUMBER: 33,771
REFERENCE/DOCKET NUMBER: 09000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 301-984-8000
TELECHART: 090158
                                                                                                                                                                                         TELEFAX: 301-230-0158
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Rockville
                                                                                                                       amino acids
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                                                                                 protein
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Score 31;
Pred. No.
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RESULT 2
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Sequence 26, Application US/08483749A
Patent No. 6034561
GENERAL INFORMATION:
APPLICANT: RING, DAVID B.
TITLE OF INVENTION: ANTIGEN-BINDIN
TITLE OF INVENTION: MOLECULES SPEC
NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                 Matches
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APPLICANT:
APPLICANT:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amind acids
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Titmas, Richard C.
APPLICANT: Williams, Richard O.
TITLE OF INVENTION: The Isolation and Production of
TITLE OF INVENTION: Catalytic Antibodies using Phage Technology
NUMBER OF SEQUENCES: 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: RYAN, JOHN W.
REGISTRATION NUMBER: 33,771
REFERENCE/DOCKET NUMBER: 09
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                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 301-984-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                          100.0%; Score 31; DB Local Similarity 100.0%; Pred. No. 6.9; 1es 7; Conservative 0. Minaria.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1530 East CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
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Chiswell, David
Darsley, Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                       92 amino acids
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Martin, Mark T.
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                                                                                                                                                                                                                                                                                                                                                                        linear
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                    ANTIGEN-BINDING SITES OF ANTIBODY MOLECULES SPECIFIC FOR CANCER ANTIGENS
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acid
            APPLICANT: Winter, Gregory
APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00013
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
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APPLICANT:
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APPLICANT:
APPLICANT:
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ZIP: 94662-8097
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: SAVEREIDE, PAUL B.
REGISTRATION NUMBER: 36,914
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 601-2585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/483,749A FILING DATE: 07-JUN-1995 CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: CHIRON CORPORATION
STREET: INTELLECTUAL PROPERTY - R440, PO BOX 8097
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FILING DATE:
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Hoogenboom, Hendricus
Griffiths, Andrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cambridge Antibody Technology Limited Medical Research Council
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                                                                                                                                                                                                                                                                                                                                                Holliger, Kasper
                                                                                                                                                                                                                                                                                                                                                                       Jackson, Ronald
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US-08-466-886-27; Sequence 27, Application US/08466886; Patent No. 5776677
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PRIOR FILING DATE: 1991-05-15
PRIOR PPLICATION NUMBER: ECT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR PPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1993-06-07
NUMBER OF SEQ ID NOS: 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. 5776677
GENERAL INFORMATION:
APPLICANT: Tsui, Land APPLICANT: Riordan,
                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/466,886
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A.
REGISTRATION NUMBER: 1329.0010006
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
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SEQ ID NO 267
LENGTH: 108
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                                                      TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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APPLICANT:
APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: Cystic Fibrosis Gene
NUMBER OF SEQUENCES: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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Local Similarity 100.0%;
nes 7; Conservative (
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ZIP: 20005
                 STRANDEDNESS:
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   COPOLOGY:
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                                      amino acid
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Colling, Francis S.
Iannuzzi, Michael C.
Drumm, Mitchell L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Riordan, John R.
Rommens, Johanna M.
                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
SS: not relevant
not relevant
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; Sequence 27, Application; Sequence 27, Application; Patent No. 6201107; GENERAL INFORMATION: APPLICANT: Tsui, Le APPLICANT: Riordan;
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                                                                                    US-08-469-617-27
                                                                                                 RESULT 7
                                                                                                                                                                                                                                                                             ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-713-939A-74
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Matches 7; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Prusiner, Stanley B.
APPLICANT: Williamson, R. Anthony
APPLICANT: Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                                                                NAME: BOZICEVÍC, KARI
REGISTRADIO NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 436
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM CON
OPERATING SYSTEM:
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ZIP: 94025
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                                                                                                                                                 50 ATSSLDS 56
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                                                                 Application US/08469617
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 Tsui, Lap-Chee
Riordan, John R.
                                                                                                                                                                                                                                                                                                                                                           109 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM Compatible
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                                                                                                                                                                                                             100.0%; Score 31; DB 2; Length 109; 100.0%; Pred. No. 8.4; Live 0; Mismatches 0; Indels
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Rommens, Johanna

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US-09-036-579-74
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,617
FILING DATE: 06-UNU-1995
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: GOldstein, Jorge A.
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 1329.0010008
                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 6290954
GENERAL INFORMATION:
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TELEPAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                  APPLICANT: Prusiner, State Applicant: Williamson, R. Anthony
APPLICANT: BY
TITLE OF INVESTIGATION ANTIBODIES SPECIFIC FOR NATIVE PrP
NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
CADDRESSE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road
STREET: 2200 Sand Hill Road
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APPLICANT:
APPLICANT:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Drumm, Mitchell L.
APPLICANT: Buckwald, Manuel
TITLE OF INVENTION: Cystic Fibrosis Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                 SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y Match 100.0%; Score 31; Di
Local Similarity 100.0%; Pred. No. 8.
hes 7; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Washington
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                                                     COMPUTER: IBM CONTROL OPERATING SYSTEM:
                                                                                                                                                                             STREET: 2200 Sand
CITY: Menlo Park
STATE: CA
                                                                                                                                                          COUNTRY:
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                                                                                                                                           94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     109 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kerem, Bat-Sheva
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                                                            IBM Compatible SYSTEM: DOS
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US/09/036,579
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Patent No. 6
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Best Local Similarity 100.0%;
Matches 7; Conservative 0
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INFORMATION FOR SEQ ID NO:
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/713
APPLICATION NUMBER: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Bozicevic, Karl
                       INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Prusiner, Stanley B.
APPLICANT: Williamson, R. Anthony
APPLICANT: Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/550
                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: 415-854-5277
                                                                                                                                                                          FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Bozicevic, Karl
                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 109 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM CONTROL SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Fish & Richardson P.C. STREET: 2200 Sand Hill Road
STRANDEDNESS:
                                                                                                                                           REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                            TELEFAX: 415-854-0875
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           amino acids
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linear
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                                                                                                                                                                                                                                                                                                                                                        IBM Compatible
single
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RESULT 11
US-08-469-630-27
JS-Sequence 27, Application US/08469630
Patent No. 6730777
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US-09-943-906-74
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                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                             Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                               TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 74:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
STRANDENNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: BOZICEVIC, Karl
REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06510.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTERQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 74:
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COMPUTER READABLE FORM:
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FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Fish & Richardson P.C. STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 415-854-0875
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ilarity 100.0%;
Conservative 0
                                                                                                                                                                                             100.0%; Score 31; DB 4; Length 109; llarity 100.0%; Pred. No. 8.4; Conservative 0; Mismatches 0; Indels
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Burton, Dennis R.
                                                                                                                          56
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Pred. No. 8.4;
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APPLICANT: Prusiner, Stanley
APPLICANT: Safar, Jiri
APPLICANT: Williamson, Anthony
APPLICANT: Williamson, Dennis
TITLE OF INVENTION: Antibodies Specific for Ungulate PrP
FILE REFERENCE: UCAL-194
CURRENT APPLICATION NUMBER: US/09/627,218B
CURRENT FILING DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO
LENGTH: 112
TYPE: PRT
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; MOLECULE TYPE: peptide
US-08-469-630-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 7
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                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/09627218B Patent No. 6537548
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                   ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: synthesized peptide
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APPLICANT:
TITLE OF IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Goldstein, Jorge A.
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/469,630 FILING DATE: 06-JUN-1995 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 109 amino acids
TYPE: amino acid
STRANDEDNESS: not relev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATION TO THE PO-POS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 202-371-2600
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                                                                                                                                                                                                                                                                                                                                                                                                                                             99 ATSSLDS 105
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Riordan, John R.
Rommens, Johanna M.
Pat-Sheva
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VENTION: Cystic Fibrosis Gene
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Iannuzzi, Michael C.
Drumm, Mitchell L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SS: not relevant
not relevant
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RESULT 13
US-08-133-804-6
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US-08-461-838-6
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Patent No.
GENERAL IN
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                                                                                         Sequence 6, Application US/08461838 Patent No. 5753204
                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                               Query Match
                                             APPLICANT: Huston APPLICANT: Opperma
                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS LENGTH: 243 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICANT: Huston, James S. APPLICANT: Oppermann, Hermann APPLICANT: Houston, L. L. APPLICANT: Ring, David B. TITLE OF INVENTION: Biosynthet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynt
TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                         100.0%;
Local Similarity 100.0%;
nes 7; Conservative
                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/133,804 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Boston
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5. 5534254
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Exchange Place, 53 State Street
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ilarity 100.0%;
Conservative (
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   Biosynthetic Binding Proteins For
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Pred. No.
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Best Local 9
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICATION NUMBER: US/08/461
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
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CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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NUMBER OF SEQUENCES:
                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                        APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynth
TITLE OF INVENTION: Imaging
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                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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amino acid
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Exchange Place, 53 State Street
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Exchange Place, 53 State Street
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                                                                                                                                                                                                                                                                                             Hurwitz & Thibeault/Patent Department
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Pred. No. 21;
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REPERENCE/DOCKET NUMBER: 2054/22

TELECOMMUNICATION INFORMATION:
TELEPHAN: 617-248-7477
TELEPHAN: 617-248-7477
TELEPHAN: 617-248-7100
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: mino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-461-386-6

Query Match
Best Local Similarity 100.0%; Score 31; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 21;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ATSSLDS 7
Db 183 ATSSLDS 189

Search completed: April 4, 2005, 16:00:52

Job time: 4.57827 secs
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Minimum
Maximum
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Maximum Match 100%
Listing first 45 summaries
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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     of hits satisfying chosen parameters:
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Match
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146.032 Million cell updates/sec
  513545 segs, 74649064 residues
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1: /cgn2_6/ptodata/1/1aa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/backfIles1.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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  US-08-483-749A-26
US-08-133-804-6
US-08-161-838-6
US-08-461-386-6
US-08-356-786-4
US-08-356-786-10
US-09-627-218B-1
US-08-273-146-45
US-08-273-146-53
US-08-888-366-20
US-08-888-366-20
US-08-888-366-26
US-09-186-219A-74
US-08-131-939A-74
US-09-155-106-21
US-09-155-106-22
US-09-155-106-24
US-09-155-106-29
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US-09-155-106-29
US-09-155-106-29
US-09-155-106-39
US-09-36-579-72
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Sequence 26, Appli
Sequence 6, Appli
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Sequence 10, Appli
Sequence 11, Appli
Sequence 13, Appli
Sequence 14, Appli
Sequence 14, Appli
Sequence 26, Appli
Sequence 27, Appli
Sequence 27, Appli
Sequence 74, Appli
Sequence 22, Appli
Sequence 23, Appli
Sequence 24, Appli
Sequence 28, Appli
Sequence 29, Appli
Sequence 30, Appli
Sequence 30, Appli
Sequence 37, Appli
Sequence 39, Appli
Sequence 72, Appli
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	Query Match Best Local Matches 1	OPERATING SYS SOFTWARE: PA CURRENT APPLICATION I ATTORNEY/AGENT NAME: SAVERI REGISTRATION REFERENCE/DON TELLEFAX: (5: INFORMATION FOR: (5: INFORMATION FOR: (5: INFORMATION FOR: (5: INFORMATION FOR: (7: SEQUENCE CHARA SEQUENCE CHARA SEQUENCE CHARA SEQUENCE TYPE: amino TOPOLOGY: 1: MOLECULE TYPE: MOLECULE TYPE:	STREET: CITY: STATE: COUNTRY ZIP: 9 COMPUTER: MEDIUM COMPUTER:	PATENT NO. 605456 GENERAL INFORMAT APPLICANT: RITTLE OF INVEN TITLE OF INVEN TITLE OF SEQUI CORRESPONDENCE ADDRESSEE:	RESULT 1 US-08-483-	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
1 RA 	cch al Simi 11;	OPERATING SY SOFTWARE: PE SOFTWARE: PE RRENT APPLICA APPLICATION PE FILING DATE: CLASSIFICATIO TORNEY/AGENT NAME: SAVERE REGISTRATION REFERENCE/DO LECOMMUNICATI TELEPHONE: (5) TELEPAX: (5) TELEFAX: (5) TELEFAX: (5) TELEFAX: 107 TYPE: amino TYPE: amino TYPE: amino TYPE: amino TYPE: dmino TYPE: dmino TYPE: dmino		INFO	4.0	
RASQDIGNSLT RASQDIGNSLT	nilarity Conserva	NUMB OTEM	E S A SE	. 6054561 INFORMATION: ANT: RING, DAVID OF INVENTION: AN OF INVENTION: MO OF OF SEQUENCES: 3 PONDENCE ADDRESS: ESSEE: CHIRON CO	it i	017713333333333333333333333333333333333
T 11	100. 100. ative	TIM: PC-DOS/MS tentIn Release TION DATA: UMBER: US/08/4 07-JUN-1995 N: 536 INFORMATION: IDE, PAUL B. INFORMATION: IDE, PAUL B. OS ON INFORMATION: 510 601-2582 EQ ID NO: 26: TERLISTICS: amino acids acid near protein	TUAL ORM: ORM:	AVID ANT MOL MOL 33 ESS: 1 COR	on US/	95 95 107 107 107 107 108 108 109 109 109 109 109 109 109 109 109 109
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	DB 3		PO B) ⁷⁰ 0		-374-72 -463-72 -463-72 -469D-1 -531B-1 -531B-1 -842A-1 -939-26 -939-26 -939-27 -245-4 -939A-73 -939A-73 -939A-73 -939A-73 -939A-73 -939A-73 -939A-73
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US-08-133-804-6

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; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-133-804-6
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                                                                                                                                                                                                                                                                                                                                    Sequence 6, Application US/08461838 Patent No. 5753204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%; Score 53; DB 1; Length 243; Best Local Similarity 100.0%; Pred. No. 0.0074; Matches 11; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                 GENERAL INFOR APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,804
FILING DATE:
CIACOTTATATA
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APPLICANT: Oppermar
APPLICANT: Houston
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TELEPHONE: 617-248-7477
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                 TITLE OF INVENTION: Biosynthetic Binding Proteins For TITLE OF INVENTION: Imaging NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Biosynthetic Binding Proteins TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 20
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                                                       COUNTRY: UZIP: 02109
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                                                                                            STATE: Massachusetts
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amino acid
                                                                                                                 Boston
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                                                                                                                           B: Testa, Hurwitz & Thibeault/Patent Department
Exchange Place, 53 State Street
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Houston, L. L.
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Houston, L. L.
Ring, David B.
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; MOLECULE TYPE:
US-08-461-386-6
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Best Local Similarity
Matches 11; Conserv
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
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                                                                                                               TELEFAX: 617-248-7100 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                   APPLICATION NUMBER: US/08/461,386
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                             REFERENCE/DOCKET NUMBER: 20.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynth
TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
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                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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                                                     amino acid
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Exchange Place, 53 State Street
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                                                                          243 amino acids
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US-08-356-786-10

; Sequence 10, Application US/08356786

; Patent No. 5877305
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                ; Patent No. 5877305; GENERAL INFORMATION:
APPLICANT: Huston,
APPLICANT: Oppermar
APPLICANT: Houston,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-356-786-4
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/356,786
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/831,967
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
NAME: Pitcher Edemind R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence 4, Application US/08356786
Patent No. 5877305
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: CR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Biosynthetic Binding Protein for Cancer TITLE OF INVENTION: Marker NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ring, David B.
                                                                                                                                                                                                                                                   Local Similarity 100.0%; es 11; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 27,829
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ilarity 100.0%;
Conservative
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Oppermann, Hermann
Houston, L. L.
Ring, David B.
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Pred. No. 0.0074;
); Mismatches 0
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APPLICANT: Prusiner, Stanley
APPLICANT: Safar, Jiri
APPLICANT: Williamson, Anthony
APPLICANT: Williamson, Anthony
APPLICANT: Burton, Dennis
TITLE OF INVENTION: Antibodies Specific for Ungulate PrP
FILE REFERENCE: UCAL-194
CURRENT APPLICATION NUMBER: US/09/627,218B
CURRENT FILING DATE: 2000-07-27
NUMBER OF SEG ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SECTUPED 11
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                                                                                   US-09-627-218B-1
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                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/09627218B Patent No. 6537548
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best
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        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 534 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/831
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                          LENGTH: 112
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                 FEATURE:
OTHER INFORMATION: synthesized peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: PItcher, Edmund R.
PItcher, Edmund R.
PItcher, Edmund R.
PREGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CRI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
B4.9%; Score 45; DB 4; Length 112; Local Similarity 90.0%; Pred. No. 0.13; 1eB 9; Conservative 1: Winner 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         448 RASQDIGNSLT 458
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Pred. No. 0.018;
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RESULT 8 US-08-273-146-45

RASODIGNNL 33 RASQDIGNSL 10

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Matches
                                                                                                    Sequence 53, App.
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                                                                           GENERAL INFORMATION:
APPLICANT: Smith,
                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 301-230-0158 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Ryan, John W.
REGISTRATION NUMBER: 33,771
REFERENCE/DOCKET NUMBER: 0900
TELECOMMUNICATION INFORMATION:
TELEPHONE: 301-984-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kentēn, John H.
APPLICANT: Martin, Mark T.
APPLICANT: Titmas, Richard C.
APPLICANT: Williams, Richard O.
TITLE OF INVENTION: The Isolation and Production of
TITLE OF INVENTION: Catalytic Antibodies using Phage Technology
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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APPLICATION NUMBER: US/
FILING DATE: 14-JUL-199
CLASSIFICATION: 435
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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                                             APPLICANT:
                  APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
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40. 5855885
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Rockville
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9; Conserv
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                                                                                                                       Application US/08273146
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Fitzgerald, Kevin
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Chiswell, David
                                            Smith, Rodger
McCafferty, John
Chiswell, David
Darsley, Michael J. Fitzgerald, Kevin Kenten, John H.
                                                                                                                                                                                                                                                             Conservative
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                                                               John
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Pred. No. 0.26;
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Best Local Similarity
Watches 9; Conservi
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Ryan, John W.
REGISTRATION NUMBER: 33,771
REFERENCE/DOCKET NUMBER: 0900
TELECOMMUNICATION INFORMATION:
TELEPHONE: 301-984-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 301-230-0158
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/273,146
FILING DATE: 14-UUL-1994
CLASSIFICATION: 435
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Martin, Mark T.
APPLICANT: Titmas, Richard C.
APPLICANT: Williams, Richard O.
TITLE OF INVENTION: The Isolation and Production of
TITLE OF INVENTION: Catalytic Antibodies using Phage Technology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/888,366
                                                                                                                           ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                               APPLICANT: Lopez, Osvaldo
APPLICANT: Wylie, Dwane E.
APPLICANT: Wagner, Fred W.
TITLE OF INVENTION: Mercur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid TOPOLOGY: linear
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                                                                                                                                                                                                                      STREET: 90 South CITY: Minneapolis
FILING DATE: 03
CLASSIFICATION:
                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 RASQDIGNSL 10
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90 South 7th Street,
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                   03-JUL-1997
                                                                                                                                                                                                                                                                                                  Mercury Binding Polypeptides and Nucleotides Coding Therefore: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  81.1%;
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Pred. No. 0.
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FILING DATE: 27-JAN-PRIOR APPLICATION DATA:

US 08/187,407 -1994 : US 07/990,542

APPLICATION NUMBER:

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US-08-888-366-14
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US-08-888-366-20
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INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
FENGTH: 107 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/888,366
PILING DATE: 03-JUL-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lopez, Osvaldo
APPLICANT: Wylie, Dwane E.
APPLICANT: Wagner, Fred W.
TITLE OF INVENTION: Mercury Binding Polypeptides and Nucleotides Coding Therefore
NUMBER OF SEQUENCES: 39
                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
FILING DATE: 27-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 35,093
REFERENCE/DOCKET NUMBER: 86.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-332-5300
               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 14-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 0
FILING DATE: 14-MAR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 14-DEC-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
PRIOR APPLICATION DATA:
                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                        COUNTRY: US
ZIP: 55402
                                                                                                                                                                                                                                                                                                                                                                      STREET: 90 South CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 81.1%;
Local Similarity 90.0%;
es 9; Conservative
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90 South 7th Street, 3100 No. 5972656west Ctr
                                                                                                                                                                                                                                                                                                                                          USA
                                                                 UMBER: US 07/990,542
14-DEC-1992
                 14-MAR-1990
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                                 US 07/493,299
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    Mismatches

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Pred. No. 0.
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RESULT 12
US-08-888-366-26
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SEQUENCE CHARACTERISTICS:
LENGTH: 107 amin --
Type
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 81.1%;
Best Local Similarity 90.0%;
Matches 9; Conservative
TELEFAX: 612-332-9081
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/
FILING DATE: 27-(JAN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 14-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Lopez, Osvaldo
APPLICANT: Wylie, Dwane E.
APPLICANT: Wagner, Fred W.
TITLE OF INVENTION: Mercury Binding Polypeptides and Nucleotides Coding Therefore
                                                                                                                                   FILING DATE: 14-MAR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                     APPLICATION NUMBER: US 0: FILING DATE: 14-MAR-1990 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 0:
                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 07/493,299
                                                                                                                                                                                                                                                                                                                                                                             SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/888,366
FILING DATE: 03-JUL-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 612-332-5300
                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 612-332-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Minneapolis
                                                                                     REGISTRATION NUMBER: 35,093 REFERENCE/DOCKET NUMBER: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Carter, Charles G. REGISTRATION NUMBER: 35, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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). 5972656
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                                                                                                                   Charles
                                                                                                                                                                                                                                                                                                                             US 08/187,407
                                                                                                                                                                         US 07/324,392
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0.31;
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; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-888-366-26
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US-09-726-219A-267
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                                                                                                                                                 , OTHER INFORMATION: light chain from clone MIF US-09-726-219A-267
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PRIOR FILING DATE: 1990-07-10
PRIOR PELICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-12
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                                                                                                                                                                                                                                                                 SEQ ID NO 267
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                         Query Match
Best Local (
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APPLICANT: Cambridge Antibody Technology
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
APPLICANT: McCafferty, John
APPLICANT: Pope, Anthony
APPLICANT: Johnson, Kevin
APPLICANT: Hoogenboom, Hendricus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 213839-00113
CURRENT APPLICATION NUMBER: US/09/726,219A
CURRENT FILING DATE: 2000-11-28
CURRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 272
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                             ENGTH: 108
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les 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/971,857
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/GB91/01134
                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 1993-01-08
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                                     1 RASQDIGNSL 10
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                                                                                             Similarity
RASODIGSSL 33
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milarity 90.0%;
Conservative
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Griffiths, Andrew
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                                                                         Conservative
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                                                                                               81.1%;
90.0%;
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L; Mismatches
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Pred. No.
                                                                                             Score 43; DB 4; Length 108; Pred. No. 0.31;
                                                                               Mismatches
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RESULT 14
US-08-713-939A-74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 74, Application US/08713939A
                                                                                                                                                                                                                                                                                                                                            Matches
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APPLICANT: Prusin
                                                                                                                                                                          Patent No. 6290954
GENERAL INFORMATION:
                                                                                                                                                                                        sequence 74, Application US/09036579 Patent No. 6290954
                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FASTSEQ VETSIC
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/O
FILING DATE: 13-SEP-1996
CLASSIFICATION: 436
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Prusiner, Stanley B.
APPLICANT: Williamson, R. Anthony
APPLICANT: Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                 APPLICANT: Prusiner, Stanley B.
APPLICANT: Williamson, R. Anthony
APPLICANT: Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE
                                                                                     CORRESPONDENCE ADDRESS
                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Bozicevic, Karl
REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
ZIP: 94025
            COUNTRY:
                                        CITY: Menlo Park
                                                       STREET:
                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                  24 RASODIGSSL
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                                                        E: Fish & Richardson P.C.
2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
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linear
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            U.S.A.
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Pred. No.
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COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: STEM: DOS
SOPTAMRE: FeatSEQ Version 2.0
CURRENT APPLICATION NUMBER: US/09/036,579
FILING DATE:
PRIOR APPLICATION NUMBER: 08/713,939
FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: BOZICEVIC, Karl
REFERNICE/DOCKET NUMBER: 06510/059001
TELECOMPUTICATION INFORMATION:
TELEPHONE: 415-854-5277
TELEFAX: 415-854-0875
TILEFAX: 415-854-0875
TILEFAX: 415-854-0875
TILEFAX: 415-854-0875
TILEFAX: 415-854-0875
TILEFAX: 415-854-0875
TILEFAX: 415-854-0875
TINDRWATION FOR SEQ ID NO. 74:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 anino acids
TYPE: anino acid
STRANDEDNESS: elngle
TODOLOGY: 11near
STRANDEDNESS: elngle
TODOLOGY: 11near
STRANDESS: peptide
US-09-036-579-74

Ouery Match
Bost Local Similarity 90.0%; Pred. NO. 0.31;
Matches 9; Conservative 1; Mismatches 0; Gaps 0;
Debt Local Similarity 90.0%; Pred. NO. 0.31;
Matches 9; Conservative 1; Mismatches 0; Gaps 0;
ARASODICSSI 33
Search completed: April 4, 2005, 16:00:51
Job time: 6.623 secs
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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ŏ.
  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length:
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Match
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2: /cgn2_6/ptodata/1/laa/5B_COMB.pep:*

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4: /cgn2_6/ptodata/1/laa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/laa/pCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/laa/backfiles1.pep:*
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US-08-461-386-6
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US-09-356-786-1
US-09-356-786-25202
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              Sequence 8, Appli
Sequence 6014, Ap
Sequence 14502, A
Sequence 3, Appli
Sequence 5104, Appl
Sequence 5104, A
Sequence 25174, A
Sequence 25045, A
Sequence 35045, A
Sequence 17424, A
Sequence 9, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
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Sequence 10, Appl
Sequence 28549, Ap
Sequence 3, Appli
Sequence 5428, Ap
Sequence 5428, Ap
Sequence 6862, Ap
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RESULT
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ALIGNMENTS	US-09-949-016-10824 US-09-949-016-10825 US-09-949-016-10826	US-09-107-433-4659 US-09-328-352-7054 US-09-489-039A-9497 US-09-902-540-10983	US-09-328-352-8007 US-09-543-681A-8072 US-09-583-110-5042	US-09-543-681A-6425 US-09-543-681A-6425 US-09-252-681A-6825 US-09-270-767-60066	-09-134 -09-583 -09-902
	Sequence 10824, A Sequence 10825, A Sequence 10826, A	Sequence 4659, Ap Sequence 7054, Ap Sequence 9497, Ap Sequence 10983, A	8007, 8072, 5042,	Sequence 12469, A Sequence 18601, A Sequence 6425, Ap Sequence 60066, A	4580, 3394, 15200,

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Sequence 24, Application US/08483749A
Patent No. 6054561
GENERAL INFORMATION:
APPLICANT: RING, DAVID B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,749A
FILING DATE: 07-UN-1995
CLASSIFICATION: 36
ATTORNEY/AGENT INFORMATION:
NAME: SAVEREIDE, PAUL B.
REGISTRATION NUMBER: 36,914
REFERENCE/DOCKET NUMBER: 05.08.008
TELLECOMMUNICATION INFORMATION:
TELLEPAX: (510) 651-2585
TELLEPAX: (510) 655-3542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (510) 655-354
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESS:
CHIRON CORPORATION
COR
                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 115 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: INTELLECTION CITY: EMERYVILLE
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REGEAY 6
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                                                                                                         Conservative
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Pred. No. 7.9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Huston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                             quence 6, Application US/08461838 tent No. 5753204
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                               APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Imaging
TITLE OF INVENTION: Imaging
                                                                                                                                             NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 617-248-7477
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Biosynthetic Binding Proteins TITLE OF INVENTION: Imaging NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                    COUNTRY: U
                                                                                                  CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 20
                                                                                                                   STREET:
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                                                                                                                                  ADDRESSEE:
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Exchange Place,
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Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 34; nilarity 100.0%; Pred. No. Conservative 0; Mismatch
                                                                    USA
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Houston, L. L.
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                                                                                                                Hurwitz & Thibeault/Patent Department Place, 53 State Street
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RESULT 4
US-08-461-386-6
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Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/08461386 Patent No. 5837846
                                                                          TELEFAX: 617-248-7100 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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APPLICATION UNIBER: US/O.
FILING DATE:
CLASSTT
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                                           SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acid
                                                                                                       REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEW PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
 MOLECULE TYPE:
                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Biosynthetic Binding Proteins For TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                 TOPOLOGY:
                                                                                                                                                    NAME: Kelley, Robin D. REGISTRATION NUMBER: 34,637
                                                                                                                                                                                                                                                                                                                                           COUNTRY: US ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Testa, Hurwit
STREET: Exchange Place,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
SOFTWARE: PatentI
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                                                                                                                                                                                                                                                                                                                                                                                            Boston
                           : 243 amino acids
amino acid
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Ring, David B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Huston, James S.
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protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hermann
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Pred. No. 16;
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RESULT 6
US-09-724-797-32
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GENERAL INFORMATION:
APPLICANT: JON S. THORSON
TITLE OF INVENTION: MICROMONOSPORA ECHINOSPORA GENES
TITLE OF INVENTION: ENCODING FOR BIOSYNTHESIS OF
TITLE OF INVENTION: CALICHEAMICIN AND SELF-RESISTANN
                                                                                          Sequence 32, Application US/09724797
Patent No. 6733998
                                                                                                                                                                                                                                                                                                                     Query Match
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Patent No. 5877305
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/831
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/356,786
                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 243 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Protein for Cancer
TITLE OF INVENTION: Marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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   AND SELF-RESISTANCE THERETO
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US-09-724-797-32
                                                                                                                                                                                                               US-08-356-786-10
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PRIOR FILING DATE: 1998-12-07
NUMBER OF SEQ ID NOS: 95
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 32
                                                                 Query Match
Best Local Similarity
Matches 6; Conserv
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APPLICANT: Huston
APPLICANT: Opperman
APPLICANT: Houston
APPLICANT: Ring, D
                                                                                                                                                                                                                                                                                                                 TELEPAX: (61/)
TELEPAX: (61/)
INFORMATION FOR SEQ ID NO: 1'
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
STORTH: 534 amino acids
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Best Local 9
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CURRENT APPLICATION NUMBER: US/09/724,797
CURRENT FILING DATE: 2000-11-28
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TYPE: PRT
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                                                                                                                                                                                                                                            MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/831,967
PILING DATE: 06-FEB-1992
PILING DATE: 140-FEB-1992
PRIOR FEBRUARY AGENT IMPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                  LENGTH: 534 amino
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Pitcher, Edmund REGISTRATION NUMBER: 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
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Similarity 100.0%;
6; Conservative 0
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   RFGFAY 6
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Houston, L. L.
Ring, David B.
                                                                        Conservative
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                                                                                                  100.0%;
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                                                                                                     Score 34; DB
Pred. No. 35;
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Pred. No. 30;
                                                                        Mismatches
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                                                                                                                                      Length 534;
                                                                        Indels
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RESULT 10
US-09-134-000C-5428
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NUMBER OF SEQ ID NOS: 15

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 3

LENGTH: 776

TYPE: PRT

ORGANIAM: S. pombe
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US-09-252-991A-28549
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                                   Sequence 5428, Application US/09134000C Patent No. 6617156
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LENGTH: 571
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APPLICANT: Marc J.
                                                                                                                                                                                                                                                                    Query Match
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BECKER, JEFFREY W.
APPLICANT: LUBKOWITZ, MARK A.
APPLICANT: LUBKOWITZ, MARK A.
TITLE OF INVENTION: ISOLATED CANDIDA ALBICANS OLIGOPEPTIDE TRANSPORTER GENE
FILE REFERENCE: 372.6520P
CURRENT APPLICATION NUMBER: US/09/165,396
CURRENT FILING DATE: 1998-10-02
EARLIER APPLICATION NUMBER: PCT/US98/02332
EARLIER FILING DATE: 1998-02-06
EARLIER APPLICATION NUMBER: 60/037,859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
APPLICANT: Lynn Doucette-Stamm et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE:
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nes 5; Conserv
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Pred. No. 3.2e+02;
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-13
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR APPLICATION NUMBER: US 60/076,409
PRIOR TILING DATE: 1998-02-13
PRIOR TILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 25202
LENGTH: 78
TYPE: PRT
                                                                         Sequence 6862, Application US/09543681A

Patent No. 6605709

GENERAL INFORMATION:
APPLICANT: GARY BRETON
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABIL)
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709,1002-001
CUCKENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 1909-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 032796-032
CURRENT APPLICATION NUMBER: US/09/134,000C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 6812
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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Best Local Similarity
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                                                              SEQ ID NO 6862
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Best Local S
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                        LENGTH: 351
TYPE: PRT
ORGANISM: Proteus mirabilis
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83.3%;
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Pred. No. 3.5e+02;
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RESULT 14
US-09-328-352-6014
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US-09-153-599A-8
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Sequence 6014, Application US/09328352
Patent No. 6562958
GENERAL INFORMATION:
APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUWANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
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Best Local 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPAX: 312-616-5460
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Weber, J. Mark
APPLICANT: Lu, B. Minh
TITLE OF INVENTION: Method for Strain Improvement of
TITLE OF INVENTION: Erythromycin Producing Bacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/153,599A
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: FE TELECOMMUNICATION INFORMATION: 312-616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Mueller, Lisa V.
REGISTRATION NUMBER: 38,978
                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
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STREET: 180 N. Stetson Avenue, 2 Prudential Plaza
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
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Pred. No. 2.7e+02;
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Pred. No. 2.4e+02;
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TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-14502
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Search completed: April Job time: 4.06709 secs

// ORGANISM: Acinetobacter baumannii
US-09-328-352-6014

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; LENGTH: 408
; TYPE: PRT
                                                                                                                                                                                                                                                                           APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 14502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14502, Application US/09248796A Patent No. 6747137
                                                                                                                                         Matches
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Similarity 100.0%; Pred. No.
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Pred. No. 3.3e+02;
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Result
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Maximum Match 100%
Listing first 45 summaries
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: /cgn2_6/ptodatta/1/1aa/5B_COMB.pep:*
3: /cgn2_6/ptodatta/1/1aa/6A_COMB.pep:*
4: /cgn2_6/ptodatta/1/1aa/6B_COMB.pep:*
5: /cgn2_6/ptodatta/1/1aa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/backfIles1.pep:*
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146.032 Million cell updates/sec
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US-08-875-811-55
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. US/08483749A ID B. ANTIGEN-BINDING SITES OF ANTIBODY MOLECULES SPECIFIC FOR CANCER ANTIGENS	ALIGNMENTS	US-09-711-485-124	US-09-610-838-124	US-09-136-389-124	US-08-839-765-124	US-08-082-842A-88	US-08-082-842A-29	US-08-082-842A-28	US-08-646-360-124	US-08-477-531B-66	US-08-477-531B-29	US-08-477-531B-28	US-08-472-788A-88	US-08-472-788A-29	US-08-472-788A-28	US-08-107-669D-66	US-08-107-669D-29	US-08-107-669D-28	US-08-477-484B-124	
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		App	App	App	App	Appl	Appl	Appl	App	Appl	App1	Appl	Appl	App1	Appl	Appl	Appl	Appl	App	

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                                                                    Query Match
Best Local S
Matches 17
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TELEPAX: (510) 655-3542
INPORMATION FOR SEQ ID NO: 24
SEQUENCE CHARACTERISTICS:
LENGTH: 115 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 33

CORRESPONDENCE ADDRESS:
ADDRESSEE: CHIRON CORPORATION
STREET: INTELLECTUAL PROPERTY - R4
CITY: EMERYVILLE
STATE: CA
COUNTRY: USA
ZIP: 94662-8097
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SORTWARE: DATEONTO Release #1 0. 1
                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION INDEER: US/08/483,749A FILING DATE: 07-JUN-1995 CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION: NAME: SAVEREIDE, PAUL B.
REGISTRATION NUMBER: 36,914 REGERENCY/DOCKET NUMBER: 36,914
                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 601-2585
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                                                                                          Similarity
                           WINTYTGQSTYADDFKE 17
WINTYTGOSTYADDFKE
                                                                      100.0%; ilarity 100.0%; Conservative 0;
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                                                                      Score 97; DB 3;
Pred. No. 2.1e-07;
0; Mismatches 0;
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US-08-133-804-6

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; TOPOLOGY: 1i
; MOLECULE TYPE:
US-08-133-804-6
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US-08-461-838-6
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APPLICANT: Huston,
APPLICANT: Oppermar
APPLICANT: Houston,
APPLICANT: Ring, Da
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Patent No. 5753204
COUNTY:
ZIP: 02109
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                 APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
RITILE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/133,804
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D
REGISTRATION NUMBER: 34,637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7100
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APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynther
TITLE OF INVENTION: Imaging
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Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                               Oppermann, Hermann
                                                                                                                                                                                                                                                                                                                    Huston, James S.
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MOLECULE TYPE: protein
US-08-461-838-6
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; MOLECULE TYPE: protein US-08-461-386-6
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                                                                                              INFORMATION FOR SEQ ID NO:
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SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                             REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Biosynthetic Binding Proteins TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Houston, L. APPLICANT: Ring, David
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,838
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                               TOPOLOGY:
                                                                                                                                                                             NAME: Kelley, Robin D. REGISTRATION NUMBER: 3
                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/461,386
                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Boston
                                           LENGTH: 243 amino acids
TYPE: amino acid
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CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                 Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Testa, Hurwitz & Thibeault/Patent Department
Exchange Place, 53 State Street
                                                                                                               617-248-7100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Huston,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; ilarity 100.0%; Conservative (
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                                                                                                                                                                                                                                 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               James S.
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                                                                                                                                                                                                                                                                                                 Version #1.25
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Best Local Similarity

Query Match

DB 2; Length 243;

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                ; Sequence 10, Applic,
patent No. 5877305
; GENERAL INFORMATIO)
; APPLICANT: Hust;
APPLICANT: Oppe;
; APPLICANT: Hous;
                                                                                                                       US-08-356-786-10
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                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 97; DB 2; Best Local Similarity 100.0%; Pred. No. 4.4e-0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/356,786
FILING DATE:
CLASSIFICATION: 424
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APPLICANT: Huston
APPLICANT: Opperm
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/8:
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynt
TITLE OF INVENTION: Marker
                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CRP-053
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
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CITY: Boston
STATE: Massacl
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                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Pitcher, Edmund R.
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                                                                   INFORMATION:
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                                                                                                     Application US/08356786
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Oppermann, Hermann
Houston, L. L.
Ring, David B.
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                                                 Huston, James S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biosynthetic Binding Protein for Cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 53 Patent No.
                                                                                                                                                                                                                                                                        Patent No. 60437...
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Rewton, Dianne L.
APPLICANT: Newton, Dianne L.
APPLICANT: Boque, Lluis
APPLICANT: Wlodawer, Alexander
APPLICANT: Wlodawer, Alexander
APPLICANT INVENTION: Recombinant Ribonuclease Proteins
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 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V6
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,811
FILING DATE: 19-FEB-1998
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TYPE: amino acide
TYPE: line-
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
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APPLICATION NUMBER: 07/831,967
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
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                                                                                                                                                                                                                STREET: Two Embarca
CITY: San Francisco
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                                                                                                                                                                                                 STATE:
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CLASSIFICATION:
                                                                                                                                                                                COUNTRY:
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17; Conserv
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                                                                                                                                                                                    USA
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100.0%; Pred. No.
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                                                                      Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9.8e-07;
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RESULT 8
US-08-875-811-55
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                                                                                                                                                                                                              ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,811
FILING DATE: 19-FEB-1998
CLASSIFICATION 435
PRIOR APPLICATION NUMBER: WO PCT/US97/02588
FRIOR APPLICATION NUMBER: WO PCT/US97/02588
FRIOR DATE: 19-FEB-1997
APPLICATION NUMBER: WO PCT/US97/02588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 55,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO:
TELEFAX: (415) 576-03 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Rybak, Susanna M.
APPLICANT: Newton, Dianne L.
APPLICANT: Boque, Lluis
APPLICANT: Wlodawer, Alexander
TITLE OF INVENTION: Recombinant Ribonuclease Proteins
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 365 amino acids
              TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                PRICE APPLICATION DATA:
APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: 011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 6
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: WO PCT/US97/02588 FILING DATE: 19-FEB-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                           NAME: Faris, Susan K.
REGISTRATION NUMBER: 41,739
REFERENCE/DOCKET NUMBER: 015280-244100US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         168
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21-FEB-1996
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93.8%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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; MOLECULE TYPE: protein US-08-875-811-55
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US-08-279-772A-6
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                                                                                                                   US-08-902-486-9
                                                                                                                                   RESULT 10
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                                                                Sequence 9, Application Patent No. 6140075
GENERAL INFORMATION:
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Best Local :
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APPLICANT: Russell, David
APPLICANT: Fuller, James
                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 608-251-9166
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27,386
REFERRNCE/DOCKET NUMBER: 11-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-251-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: United States of America
ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICANT: RUSSEL, DAVID R.
APPLICANT: FULLEY, JAMES T.
TITLE OF INVENTION: METHOD FOR PRODUCING ANTIBODIES AND
TITLE OF INVENTION: PROTEIN TOXINS IN PLANT CELLS
                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 252 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                      TYPE:
                                                                                                                                                                                                                                                   Local Similarity 93.8
les 15; Conservative
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es 15; Conserv
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                                                                                                                                                                                   180 WINTYTGOPTYADDFK 195
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                                                                                                  Application US/08902486
                                                                                                                                                                                                                                                                                                                                                                                   252 amino acids
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                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                     89.7%;
93.8%;
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Pred. No. 1.1e-05;
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Pred. No. 1
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NUMBER OF SEQUENCES: 1

CITY: Madison STREET:

ADDRESSEE: Quarles & Brady STREET: 1 South Pinckney Street

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APPLICANT: Buyes, Marie-Ange
APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISBASES AND SKIN DISORDERS
FILE REFERENCE: INN8:015
CURRENT APPLICATION NUMBER: US/09/485,737B
CURRENT FILING DATE: 2000-02-14
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-06-14
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR APPLICATION NUMBER: EPO 98870122.5
PRIOR FILING DATE: 1998-06-18
OFTWARE: PATE: 1997-08-18
NUMBER OF SEQ ID NOS: 104
SOFTWARE: Patentin version 3.0
SEQ ID NO 102
LENGTH: 230
TYPE: PRT
                                                                                     ; ORGANISM: Artificial Sequence; PEATURE; OTHER INFORMATION: SYNTHETIC US-09-485-737B-102
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/ MOLECULE TYPE: protein
US-08-902-486-9
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INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
LENGTH: 252 amino acids
TYPE: amino acid
TYPE: amino acid
                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 102, Application US/09485737B Patent No. 6350860
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Best Local Similarity
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      Matches
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PILING DATE:
CLASSIFEICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 670
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ZIP: 53701-2113
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 608-251-5000
/ Match 87.6%; Score 85; DB 3; Length 230
Local Similarity 87.5%; Pred. No. 2.7e-05;
les 14; Conservative 1; Mismatches 1; Indels
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llarity 93.8%; Pred. No. 1.
Conservative 0; Mismatche
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                                         Length 230;
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    Gaps
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APPLICANT: Sabion, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 09/485,737
PRIOR FILING DATE: 1998-08-14
PRIOR FILING DATE: 1998-08-14
PRIOR PRIOR PILING DATE: 1998-06-18
PRIOR PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR PILING DATE: 1998-06-18
PRIOR PILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
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US-10-071-485-102
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US-09-485-737B-93
                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/485,7378
CURRENT FILING DATE: 2000-00-14
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-08-14
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1998-06-18
PRIOR PPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
NUMBER OF SEQ ID NOS: 104
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                                                                                                                SEQ ID NO 93
LENGTH: 235
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 93, Application US/09485737B Patent No. 6350860
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LENGTH: 230
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Buyse, Marie-Ange
APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERPERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.0
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APPLICANT: Sablon, Erwin
                         OTHER INFORMATION: SYNTHETIC
                                                        ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ch 87.6%;
l Similarity 87.5%;
l4; Conservative
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; Pred. No. 2.7e:
1; Mismatches
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APPLICANT: Buyse, Marie-Ange
APPLICANT: Sablon, Erwin
ITILE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
ITILE OF INVENTION: SHOCK,
ITILE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
IF LE REFERENCE: INNS:015
CURRENT APPLICATION UNMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 09/485,737
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: EPCO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR FILING DATE: 1997-08-18
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US-09-485-737B-91
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Best Local Similarity 87.5
"Arches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 93, Applica Patent No. 6830752 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 104
SOFTWARE: PatentIn version 3.0
SEQ ID NO 93
LENGTH: 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 91, Applica Patent No. 6350860 GENERAL INFORMATION:
                                                                       SOFTWARE: PatentIn version 3.0 SEQ ID NO 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 87.6%; Score 85; DB 4; Best Local Similarity 87.5%; Pred. No. 2.8e-05; Matches 14; Conservative 1; Mismatches 1
                                                                                                              APPLICANT: Buyse, Marie-Ange
APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFEBRON-gamma-BINDING MOLECULES FOR TREATING SEPTIC SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
CURRENT FILLING DATE: 2000-02-14
CURRENT FILLING DATE: 2000-02-14
PRIOR APPLICATION NUMBER: ECT/EP 98/05165
PRIOR FILLING DATE: 1998-08-18
PRIOR FILLING DATE: 1998-08-18
PRIOR FILLING DATE: 1998-08-18
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1999-08-18
PRIOR FILING DATE: 1999-08-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
                       LENGTH: 24
TYPE: PRT
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ORGANISM: Artificial Sequence
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Pred. No. 2.8e-05;
1; Mismatches 1.
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; FEATURE:
; OTHER INFORMATION: SYNTHETIC
US-09-485-737B-91

Query Match
Best Local Similarity 87.5%; Pred. No. 2.8e-05;
Best Local Similarity 87.5%; Pred. No. 1.8e-05;
Matches 14; Conservative 1; Mismatches 1; Indels 0; Gaps
Qy 1 MINTYTGOSTYADDFK 16
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Db 50 WINTYTGESTYVDDFK 65

Search completed: April 4, 2005, 16:00:50
Job time: 8.6901 secs
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Sequence

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No.
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Maximum Match 100%
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Perfect score:
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US-08-783-853A-84

US-09-344-050-94

US-09-344-050-20

US-08-483-749A-24

US-08-483-749A-24

US-08-483-66-66

US-08-483-749A-66

US-08-485-336-124

US-08-485-336-124

US-08-488-138-126

US-08-488-138-126

US-08-477-484B-126

US-08-107-669D-29

US-08-107-669D-29

US-08-472-788A-89

US-08-472-788A-89

US-08-472-788A-89
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127, App

128, Appl

29, Appl

67, Appl

68, Appl

89, Appl

89, Appl
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US-08-783-853A-8
; Sequence 8, Applicatio
; Patent No. 6005091
; GENERAL INFORMATION:
                                                                                                                                        APPLICATION UMBER: US/08/783,853A
FILING DATE: 16-JAN-1997
CCLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/029,119
FILING DATE: 24-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Baumeister, Kirk
REGISTRATION NUMBER: 3,833
REGISTRATION NUMBER: 3,833
                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 19406
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: LEM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Blackburn, Michael
APPLICANT: Church, William
APPLICANT: Gross, Mitchell
APPLICANT: Feuerstein, Giora
APPLICANT: Nichols, Andrew
APPLICANT: Nichols, Andrew
APPLICANT: Padlan, Eduardo
APPLICANT: Patel, Arunbhai
APPLICANT: Sylvester, Daniel
TITLE OF INVENTION: ANTICOAGULANT ACTITLE OF INVENTION: OF THROMBOSIS
NUMBER OF SEQUENCES: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: P5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5096
  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
            TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: SmithKline Beecham Corporation STREET: 709 Swedeland Road CITY: King of Prussia STATE: PA
                                               LENGTH:
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US-08-646-360-124

US-08-646-360-126

US-08-646-360-126

US-08-082-842A-29

US-08-082-842A-89

US-08-082-842A-89

US-08-083-765-124

US-08-083-765-126

US-09-136-389-126

US-09-136-389-126

US-09-610-838-124

US-09-610-838-124

US-09-610-838-126

US-09-410-781-96

US-09-711-485-126
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; FRAGMENT TYPE: ; ORIGINAL SOURCE: US-09-344-050-8
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US-09-344-050-8
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 19406

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS

SOFTWARE: FASTSEQ Version 1.5

CURRENT APPLICATION DATA:

APPLICATION UMBER: US/09/344,05

FILING DATE: 24-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/78:
APPLING DATE: 16-JAN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL: NO ANTI-SENSE: NO FRAGMENT TYPE: i ORIGINAL SOURCE:
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                                                                                                         HYPOTHETICAL:
                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 33,833
REFERENCE/DOCKET NUMBER: P5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Padlan, Eduardo
APPLICANT: Patel, Arunbhai
APPLICANT: Sylvester, Daniel
TITLE OF INVENTION: ANTICOAGULANT AGENTS USEFUL IN TREATMENT
TITLE OF INVENTION: OF THROMBOSIS
NUMBER OF SEQUENCES: 111
                                                                                                ANTI-SENSE:
                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                            TOPOLOGY:
                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: PA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: King of Prussia
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                                                                                                                                                                                         ENGTH:
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Church, William
Gross, Mitchell
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    100.0%;
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    Score 30; DB 3;
Pred. No. 4.1e+05;
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US-08-783-853A-84
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Best Local Similarity
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ITILE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
ITILE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 3503
LENGTH: 74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tent No.
                                                                                                                                                    ZIP: 19406
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0:
FILING DATE: 24-OCT-1996
                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PastESQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                                            APPLICATION NUMBER: US/0 FILING DATE: 16-JAN-1997 CLASSIFICATION:
                                                                                                                                                                                                                           ADDRESSEE: SmithKline Beecham Corporation STREET: 709 Swedeland Road CITY: King of Prussia STATE: PA
                                                                                                                                                                                              COUNTRY: UZIP: 19406
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08783853A
                                                                                                                                                                                                                                                                                                                                                                                                  Church, William Gross, Mitchell Feuerstein, Giora Nichols, Andrew
                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                         Patel, Arunbhai
Sylvester, Daniel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                               Diskette
                                                                                                                                                                                                                                                                                                                            ANTICOAGULANT
OF THROMBOSIS
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               60/029,119
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Pred. No. 24;
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                                                                                                                                                                                                                                                                                                                                           AGENTS USEFUL IN TREATMENT
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ATTORNEY/AGENT INFORMATION:

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US-09-344-050-84
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FRASLEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/344,05;
FILING DATE: 24-JUN-1999
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/783,853
FILING DATE: 16-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Baumeister, Kirk
REGISTRATION NUMBER: 33,833
REFERENCE/DOCKET NUMBER: P50438
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Fewerstein, Giora
APPLICANT: Nichols, Andrew
APPLICANT: Padlan, Edwardo
APPLICANT: Padlan, Edwardo
APPLICANT: Sylvester, Daniel
APPLICANT: Sylvester, Daniel
TITLE OF INVENTION: ANTICOAGULANT AGENTS USEFUL IN TREATWENT
TITLE OF INVENTION: OF THROMBOSIS
NUMBER OF SEQUENCES: 111
                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
ANTI-SENSE: 1
                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 610-270-5096
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Local Similarity 100.0%;
hes 5; Conservative (
                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
ZIP: 19406
                                                                                                                                                                                                                                                                                                                                                                         STREET: 709 Swedeland CITY: King of Prussia STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Baumeister, Kirk REGISTRATION NUMBER: 33,833 REFERENCE/DOCKET NUMBER: P50438
TELEX:
               TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Blackburn, Michael
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                                                                 P50438
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Pred. No. 30;
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RESULT 6
US-08-783-853A-20
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                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFO
                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/783,8
FILING DATE: 16-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: p
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: i:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                    ZIP: 19406
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                              NAME: Baumeister, Kirk
REGISTRATION NUMBER: 33,833
REFERENCE/DOCKET NUMBER: P50438
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5096
                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/029,119
FILING DATE: 24-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
CITY: King of Prussia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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MOLECULE TYPE:
                                                                                                                  TELEFAX:
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                             STRANDEDNESS:
                                           TYPE: amino acid
                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                 TOPOLOGY:
                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                              T: Church, Will.
T: Gross, Mitchell
NT: Feuerstein, Giora
NT: Nichols, Andrew
ANT: Padlan, Eduardo
ANT: Patel, Arunbhai
CANT: Sylvester, Daniel
SOF INVENTION: OF THROMBOSIS
F. OF INVENTION: OF THROMBOSIS
TOTHRICES: 111
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6005091
                                                          112 amino acids
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Church, William
               linear
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; FRAGMENT TYPE:
; ORIGINAL SOURCE:
US-09-344-050-20
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US-09-344-050-20
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Query Match 100.0%; Score 30; DB 3; Length 112; Best Local Similarity 100.0%; Pred. No. 36; Matches 5; Conservative 0; Mismatches 0; Indels
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Sequence 20, Appr-
No. 6391299
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                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/09/344,050
FILING DATE: 24-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Blackb
APPLICANT: Church
                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: Baumeister, Kirk
REGISTRATION NUMBER: 33,83
REFERENCE/DOCKET NUMBER: P504
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5096
                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 112 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO FRAGMENT TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/7
FILING DATE: 16-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Nichols, Andrew
APPLICANT: Paddan, Eduardo
APPLICANT: Patel, Arunbhai
APPLICANT: Sylvester, Daniel
TITLE OF INVENTION: ANTICOAGULANT AGENTS USEFUL IN TREATMENT
TITLE OF INVENTION: OF THROMBOSIS
NUMBER OF SEQUENCES: 111
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                       ANTI-SENSE:
                                                                                                                   MOLECULE TYPE:
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                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 19406
                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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                                                                                                                                                                                          112 amino acids
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                                                                                                       NO
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Gross, Mitchell
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                                                                                                                                                  linear
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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 Gaps
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US-08-483-749A-24
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US-08-249-013-6
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                              ZIP: 90067
                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                         COUNTRY:
                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                             USA
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Sequence 6, Application US/08249013
Patent No. 5643754
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STAIL.

COUNTRY: USA
ZIP: 94662-8097

ZIP: 94662-8097

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPER: Floppy disk
PC-DOS/MS-DOS
Palease #1.0,
                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Haake, David A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (510) 655-354
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 115 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: RING, DAVID B.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: SAVEREIDE, PAUL B.
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
TO (100/100 7400
                                                                                                                                                                 NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 601-2585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                        CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995 CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity es 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                          31 NYGMN 35
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6054561
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INTELLECTUAL PROPERTY - R440, PO BOX 8097
                                                                                                                                 E: Spensley Horn Jubas & Lubitz
1880 Century Park East, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              115 amino acids
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ilarity 100.0%;
Conservative 0
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79. 36,914
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                                                                                                                                                                                                      CLONED leptospira OUTER MEMBRANE PROTEIN
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OPERATING SYSTEM: PC-DOS/MS-DOS

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US-08-886-863-6
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                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/249,013
PILING DATE: 25-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Tumarkin Ph.D., Liss A.,
REGISTRATION NUMBER: P-38,347
REFERENCE/DOCKET NUMBER: PD-3602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Haake, David A.
TITLE OF INVENTION: CLONED
                                  TELEFAX: (619) 455-5110 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (619) 455-5110 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: FORM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/886,863
FILING DATE: 01-UU1-1997
CLASSIFICATION: 435
CLASSIFICATION: 435
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LENGTH: 117 amino acids
SEQUENCE CHARACTERISTICS:
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NAME: Tumarkin Ph.D., Lisa A.,
REGISTRATION NUMBER: P-38,347
REFERENCE/DOCKET NUMBER: PD-3602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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CITY: Los Angeles
STATE: California
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                                                                                                                        TELEPHONE:
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1880 Century Park East, Suite 500
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1..117
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                                                                                                                        (619) 455-5100
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; NAME/KEY: Protein
; LOCATION: 1..117
US-09-175-229-6
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Best Local Similarity
Watches 5; Conserve
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                            Query Match 100.0%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 38; Matches 5; Conservative 0; Mismatches
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Patent No. 6309641
                                                                                                                                                                                                                                                                            TELEFAX: (619) 455-511 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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TITLE OF INVENTION: CLONED leptospira OUTER MEMBRANE PROTEIN
NUMBER OF SEQUENCES: 10
                                                                                                                                                                   IMMEDIATE SOURCE:
                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 455-5100
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 25-MAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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LOCATION:
                                                                                                                                                                                                                                                                                                                                     NAME: Tumarkin Ph.D., Li
REGISTRATION NUMBER: P-3
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Spensley Horn Jubas & Lubitz
STREET: 1880 Century Park East, Suite 500
CITY: Los Angeles
STATE: California
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STRANDEDNESS: si
                                                                                                                                                      CLONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                 TOPOLOGY:
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1 NYGMN 5
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Tumarkin Ph.D., Lisa ....
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                                                                                                                                                                                                                                                                                                                                                                     Lisa A.,
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                                                             DB 3;
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                                                            Length 117;
                                 Indels
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PCT-US95-06764-6

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RESULT 13
US-08-425-336-124
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                                                                                                                                                                                                 Sequence 124, Application US/08425336 Patent No. 5621083
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 5; Conserv
                                                                                      GENERAL INFORMATION:
APPLICANT: Better, Marc D.
APPLICANT: Carroll, Stephen F.
APPLICANT: Studnika, Gary M.
TITLE OF INVENTION: Immunotoxir
TITLE OF INVENTION: Proceins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: The Regents of the University of California TITLE OF INVENTION: CLONED leptospira OUTER MEMBRANE PROTEIN NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Haile, Ph.D., Lisa A.,
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: FD3602
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 678-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: line MOLECULE TYPE: p IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: PCT/US95/06764
FILING DATE: 25-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                     CORRESPONDENCE ADDRESS:
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                                                                           NUMBER OF SEQUENCES:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
     STREET:
CITY: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: si
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                                      ADDRESSEE:
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                    6300 Sears Tower,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               117 amino acids
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                                                                                                           Immunotoxins Comprising Ribsome-Inactivating
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                    O'Toole, Gerstein, Murray & Ower, 233 South Wacker Drive
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                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1:
Patent No.
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Best Local Similarity 100.
Matches 5; Conservative
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TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                          STREET: Chicago
CITY: Chicago
TTATE: Illinois
TISA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Meyers, Thomas C.
REGISTRATION NUMBER: P-36,989
REFERENCE/DOCKET NUMBER: 31394
TELECOMMUNICATION: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/787,567
FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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FILING DATE: 19-JUN-1992
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                      APPLICANT: Better, Marc D.
APPLICANT: Carroll, Stephen F.
APPLICANT: Studnika, Gary M.
TITLE OF INVENTION: Immunotoxins Comprising Ribsome-Inactivating
TITLE OF INVENTION: Proteins
                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMBER: FILING DATE: 18-APP
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 18-APR-1995
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                      US/08/425,336
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APPLICATION NUMBER: 08/064,691 FILING DATE: 12-MAY-1993

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RESULT 15
US-08-488-113B-124
, Sequence 124, Application US/08488113B
, Patent No. 5744580
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APPLIANCE
PILING DATE: 12-MAY-1993
APPLICATION NUMBER: US 07/901,707
FILING DATE: 19-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/787,567
FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
AMAE: Meyers, Thomas C.
REGISTRATION NUMBER: P-36,989
REFERENCE/DOCKET NUMBER: 31394
TRLECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
TELEPAX: 312/474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPAX: 312/474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 126:
SEQUENCE CHARACTERISTICS:
LENGTH: 118 amino acids
TYPE: amino acid
                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/488,113B
FILING DATE: 07-JUN-1995
CLASSIFICATION TO BATA:
APPLICATION NUMBER: US 08/425,336
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/64,691
FILING DATE: 18-APPLICATION DATA:
APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/988,430
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/988,430
FILING DATE: 09-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/988,430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 30; DB 1; Length 118; Best Local Similarity 100.0%; Pred. No. 38; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Better, Marc D.
APPLICANT: Carroll, Stephen F.
APPLICANT: Studnika, Gary M.
TITLE OF INVENTION: Immunotoxins Comprising Ribosome-Inactivating
TITLE OF INVENTION: Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 500 West
CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U
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STRANDEDNESS: 81.
APPLICATION NUMBER: US 07/901,707 FILING DATE: 19-JUN-1992
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500 West Madison Street, 34th floor
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Search completed: April Job time: 3.55591 secs
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Best Local Similarity luv.
Watches 5; Conservative
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TELEX: 650 388-1248
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                              LENGTH: 118 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          NAME: McNicholas, Janet M. REGISTRATION NUMBER: 32,918 REFERENCE/DOCKET NUMBER: 11
                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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                                                             31 NYGMN 35
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Result
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Perfect score:
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               Pred. No. is the score greater tand is derived
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is derived by analysis of the total score distribution.
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length: 2000000000
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Match
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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Oy 1 EIQLYQSGPELKKPGETYKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY	Query Match 100.0%; Score 1258; DB 1; Best Local Similarity 100.0%; Pred. No. 3.9e-105 Matches 243; Conservative 0; Mismatches 0;	INFORMATION FO SEQUENCE CHA LENGTH: 2 TYPE: ami TOPOLOGY: MOLECULE TYP S-08-133-804-6	ATION NUCE CE/DOCKE CE/DOCKE NICATION NE: 617-2	APPLICATION NUMBER: US/08/133,804 FILING DATE: CLASSIFICATION: 424 ATTORNEY/AGENT INFORMATION: NAME: Kelley Robin D.	OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.2 CURRENT APPLICATION DATA:	ZIP: 02109 COMPUTER READABLE F MEDIUM TYPE: F10	CITY: STATE: COUNTR	CORRESPONDENCE ADDRESS: ADDRESSEE: Testa, Hurwitz & Thibeault/Patent STREET: Exchange Place, 53 State Street	TITLE OF INVENTION: BIOSYNTHETIC BINDING PROCES TITLE OF INVENTION: Inaging NUMBER OF SEQUENCES: 11	APPLICANT: Oppermann, Hermann APPLICANT: Houston, L. L. APPLICANT: Ring, David B.	Patent No. 5534254 GENERAL INFORMATION APPLICANT: Husto	ESULT 1 S-08-133-804-6	ALIGNMENTS	4 779 61.9 235 5 766.5 60.9 230	82.5 62.2 287 4 US-09-318-786- 779 61.9 235 3 US-09-485-737E	9 799.5 63.6 366 3 US-08-875-811- 791.5 62.9 240 3 US-09-485-737E	6 803 63.8 284 3 US-09 7 803 63.8 284 4 US-09 8 799.5 63.6 365 3 US-08	4 804 63.9 711 3 US-09-485-737E 5 804 63.9 711 4 US-10-071-485-	2 804 63.9 541 3 US-C 3 804 63.9 541 4 US-1	0 804 63.9 267 3 US-09-485-737B-2	8 818.5 65.1 241 2 US-08-465-473B-
CKASGYTFANYGWNWMKQAPGKGLKWMGWINTYTGQSTY 60 	58; DB 1 . 3.9e-10 ches 0			۵	Version #1.			/Pat et	Prot				NMENTS	1-485-93 Sequence 93, App 5-737B-102 Sequence 102, Ap	Sequence 37, Sequence 37,	Sequence 55, App Sequence 91, App	sequence 40, App Sequence 40, App Sequence 53, App	Sequence 90, App Sequence 90, App	Sequence 85, App	Sequence 2,	Sequence 5, Appl

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Matches 243; Conserv
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
LENGTH: 243 amino aci
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APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 20 TELECOMMUNICATION INFORMATION: TELEPHONE: 617-248-7477
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                     TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                         NAME: Kelley, Robin D. REGISTRATION NUMBER: 34,637
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
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ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS
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Exchange Place, 53 State Street
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                                                                                                                                  100.0%; Score 1258; DB 1; ilarity 100.0%; Pred. No. 3.9e-105; Conservative 0; Mismatches 0;
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TOPOLOGY: linear
; MOLECULE TYPE: protein
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Patent No.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENCTH: 243 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,386
FILING DATE:
CLASSIFICATION: 424
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NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 201
TELECOMMUNICATION INFORMATION:
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APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
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MEDIUM TYPE: Floppy disk
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STATE: Massachusetts
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TELEFAX: 617-248-7100
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5. 5837846
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                                                                                                                                                                                1 EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
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IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLEIK 240
                                                    SGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRL 180
                                                                                               ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS
                                                                                                                 ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS
                                                                                                                                                               EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
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Exchange Place,
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Place, 53 State Street
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Pred. No. 3.9e-105;
); Mismatches 0;
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APPLICANT: Huston, James S.
APPLICANT: Oppermann, Herman.
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynth
TITLE OF INVENTION: Marker
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 1258; DB 2; Best Local Similarity 100.0%; Pred. No. 3.9e-105; Matches 243; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
TYPE: amino acid
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/08356786 Patent No. 5877305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/831,967
FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/356,786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault
STREET: Exchange Place, 53 State Street
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MEDIUM TYPE: Ploppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Pitcher, Edmund R.
REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CR
 181
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                     IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLEIK 240
                                                                                                  SGSSSSGSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRL 180
                                                                                                                                                  ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQG'TLVSVSASISSS
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                                                                                                                                                                                                                                                                                                                                        Length 243;
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US-08-356-786-10
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Patent No.
                                                                                                                                                                                                                                                                                                                                          Matches 235; Conservative
                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Pitcher, Edmund R.
REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CRP-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEPAX: (617) 248-7100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (617) 248-710
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 07/831
PILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/356,786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Huston, James APPLICANT: Oppermann, Hern APPLICANT: Houston, L. L. APPLICANT: Ring, David B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
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ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz, & Thibeault
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Biosynthetic Binding Protein TITLE OF INVENTION: Marker
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: ami
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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IKRAD 243
                                     RLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLE
                                                           RLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLE
                                                                                                            GGGGGGGGGGGGDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLSQEPDGTIK
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APPLICANT: Huston,
APPLICANT: Oppermar
APPLICANT: Houston,
APPLICANT: Ring, Da
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,804
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
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5. 5534254
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                                                                                                                                                                                                                  ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRF----GFAYWGQGTLVSVSAS 116
LEIKRAD 245
                               LEIKRAD 243
                                                            PKLLIYWTSTRHTGVPDRFTGSGSGTDYTLTISSVQAEDLALHYCQQHYRVPYTFGGGTK 238
                                                                                             IKRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTN 236
                                                                                                                                                                                             AEEFKGRFAFSLETSASTAYLQINNLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA- 121
                                                                                                                            -SSSSGSSSSGSSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAWYQQKPGQS 178
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Houston, L. L.
Ring, David B.
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71.7%;
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                                                                                                                                                                                                                                                                                                                             Score 914; DB 1;
Pred. No. 3.1e-74;
02; Mismatches 30
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                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 250;
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RESULT 7

RESULT 8
US-08-463-675-8
; Sequence 8, Application US/08463675
; Patent No. 5658763

GENERAL INFORMATION:

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                                                                                                                                                                                                                                                                                                                                                                  Matches 177; Conservative
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TELEFAX: 617/248-7100
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: KELLEY, ROBIN D
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRPO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/248-7000
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, INC
STREET: 45 SOUTH STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: DORAI, HAII APPLICANT: OPPERMANN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435
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                                LEIKRAD 243
                                                                      PKLLIYWTSTRHTGVPDRFTGSGSGTDYTLTISSVQAEDLALHYCQQHYRVPYTFGGGTK 238
                                                                                                                                                                                                                    ABEFKGRFAFSLETSASTAYLQINNLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA-
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LEIKRAD 245
                                                                                             IKRLIYATSSIDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTN 236
                                                                                                                                           -SSSGSSSGSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAMYQQKPGQS
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RESULT 9
US-08-464-589-8
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Sequence 8, Application US/08464589
PATENT NO. 5733782
PATENT NO. 5733782
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: DORAI, HAIMANTI
APPLICANT: OPERMANN, HERMANN
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HIGH PROTEIN
TITLE OF INVENTION: PRODUCTION FROM NON-NATIVE DNA
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Best Local Similarity
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FILING DATE: 05-JUN-1995
CLASSIFICATION: 435
CRISTICATION DATA:
APPLICATION NUMBER: 08/143,498
FILING DATE: 25-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: KELLEY, ROBIN D
REGISTRATION NUMBER: 34,637
REPERENCE/DOCKET NUMBER: CRP093
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/248-7000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: DORAI, HAIMANY:
APPLICANT: OPPERMANN, HERMANN
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HIGH PROTEIN
TITLE OF INVENTION: PRODUCTION FROM NON-NATIVE DNA
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, INC
STREET: 45 SOUTH STREET
CITY: HOPKINTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 250 amino acids
TYPE: amino acid
TOPOLOGY: linear
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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RESULT 10
US-08-461-838-2
; Sequence 2, Application US/08461838
; Patent No. 5753204
; GENERAL INFORMATION:
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; LENGTH: 250 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-464-589-8
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FILING DATE: 25-OCT-1993
ATTORNEY, AGENT INFORMATION:
NAME: KELLEY, ROBIN D
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRP093
TELECOMMUNICATION INFORMATION:
TELEPAN: 617/248-7100
TELEPAN: 617/248-7100
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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CORRESPONDENCE ADDRESS:
ADDRESSEE: PATENT ADM.
STREET: 45 SOUTH STREE
CITY: HOPKINTON
STATE: MA
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Houston, L. L.
TITLE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/464,589 FILING DATE: 05-JUN-1995 CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABEFKGRFAFSLETSASTAYLQINNLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, INC. 45 SOUTH STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              72.7%; Score 914; DB 1; Length 250; ilarity 71.7%; Pred. No. 3.1e-74; Conservative 32; Mismatches 30; Indels
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RESULT 11
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                                                                                                                                                                                                                                                           Sequence 2, Application US/08461386 Patent No. 5837846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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COMPUTER: IBM PC COMPARIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/461,838
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                              TITLE OF INVENTION: Biosynthetic Binding Proteins TITLE OF INVENTION: Imaging NUMBER OF SEQUENCES: 11
                                                                                                                                                      APPLICANT: Huston, James S.
APPLICANT: Oppermann, Herma
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                         CORRESPONDENCE ADDRESS:
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CITY: B
                                       STREET:
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                                                        ADDRESSEE:
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                Boston
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Massachusetts
                         E: Testa, Hurwitz & Inipeauit/F
Exchange Place, 53 State Street
                                                                                                                                                      Oppermann, Hermann
Houston, L. L.
Ring, David B.
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                                                      Hurwitz & Thibeault/Patent Department
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   72.7%; Score 914; DB 1; Length 250; 71.7%; Pred. No. 3.1e-74; tive 32; Mismatches 30; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                 US-08-356-786-16
                                                                                                                                                                                                    Patent No. 5877305
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Protein
TITLE OF INVENTION: Marker
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                                                           Sequence 16,
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 250 amino acid
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Edmund R. Pitcher, Testa, Hurwitz,
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 209
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,386
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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TYPE: amino acid
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CLASSIFICATION: 424
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5877305
                                                                                                                                Boston
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                                                                                                              Massachusetts
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US-08-488-113B-148
; Sequence 148, Application US/08488113B
; Patent No. 5744580
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PATENT NO. 5744580

GENERAL INFORMATION:

APPLICANT: Better, Marc D.

APPLICANT: Carroll, Stephen F.

APPLICANT: Studnika, Gary M.

APPLICANT: Studnika, Gary M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 622 amino acid
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REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CRI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: UPILING DATE: CLASSIFICATION: 424 PRIOR APPLICATION DATA:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                      STREET: 500 West
CITY: Chicago
STATE: Illinois
                                                                                                                    COUNTRY: U
                                                                                                                                                                                                                     ADDRESSEE:
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amino acid
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                                                                                                                                                                                                 500 West Madison Street, 34th floor
                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (617)
                                                                                                                                                                                                                   McAndrews, Held & Malloy, Ltd.
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                                                                                                                                                                                                                                                                            Proteins
                                                                                                                                                                                                                                                                                             Immunotoxins Comprising Ribosome-Inactivating
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                                                          Patent No. 5756699
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Better, Marc D.
APPLICANT: Carroll, Stephen F.
APPLICANT: Studnika, Gary M.
TITLE OF INVENTION: Immunotoxins Comprising Ribosome-Inactivating
TITLE OF INVENTION: Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 312/707-8889
TELEFAX: 312/707-9155
TELEX: 650 388-1248
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 240 amino acids
                                                                                                                                                                                      Sequence 148, Application US/08477484B Patent No. 5756699
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                                   NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 312/707-8889
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 19-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
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PRIOR APPLICATION NUMBER: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 32,918
REFERENCE/DOCKET NUMBER: 11022US07/200-70.P3.C2A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: MCNICHOLAS, UN REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 0: FILING DATE: 18-APR-1995
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               ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SSSSSSSSSSSSSSSSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTI 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARR---FGFAYWGQGTLVSVSASI 117
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E: McAndrews, Held & Malloy, Ltd. 500 West Madison Street, 34th flo
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117

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STREET:

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com

Illinois

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US-08-646-360-148; Sequence 148, Application US/08646360; Patent No. 5837491; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION UNDER: US 07/988,430
FILING DATE: 09-DEC-1992
PRIOR APPLICATION UNDER: US 07/901,707
APPLICATION NUMBER: US 07/901,707
FILING DATE: 19-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/787,567
APPLICATION NUMBER: US 07/787,567
FILING DATE: 04-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 650 388-1248
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 0 FILING DATE: 04-NOV-1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 18-APR-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PILING DATE: 07-JUN-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/425,336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,484B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: McNicholas, Janet M.
REGISTRATION NUMBER: 32,918
REFERENCE/DOCKET NUMBER: 11022US07/200-70.P3.C2A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEFAX: 312/707-9155
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65.8%; Pred. No. 1.9e-66;
Live 28; Mismatches 49
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 240 amin
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Best Local :
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
FILING DATE: 12-WAY-1993
                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
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APPLICATION NUMBER: 1
FILING DATE: 19-JUN-
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 312/707-900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: McNicholas, Janet M. REGISTRATION NUMBER: 32,91 REFERENCE/DOCKET NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/01 FILING DATE: 13-MAY-1996
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 0 FILING DATE: 04-NOV-1991
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178
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                                                                                                                                                                 ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARR---FGFAYWGQGTLVSVSASI 117
                                                                                                                                                                                                EIQLVQSGGGLVKPGGSVRISCAASGYTFTNYGMNWVRQAPGKGLEWMGWINTHTGEPTY
KTLIYRANRLESGVPSRPSGSGSGTDYTLTISSLQYEDFGIYYCQQYDESPWTFGGGTKL
                             KRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNL
                                                                                             SSSSGSSSSGSSSGSDIOMTOSPSSLSASLGERVSLTCRASODIGNSLTWLOOEPDGTI
                                                                  SGGGGSGGGGGGSDIQMTQSPSSLSASVGDRVTITCRASQDINSYLSWFQQKPGKAP
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Carroll, Stephen I
Studnika, Gary M.
                                                                                                                                                                                                                                                                      Conservative
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                    65.7%; Score 827; DB 2; 65.8%; Pred. No. 1.9e-66;
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Minimum
Maximum
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Perfect score:
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Gapop 10.0 ,
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53
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193.690 Million cell updates/sec
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1: geneseqp1980s:*
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Maximum Match 100%
Listing first 45 summaries
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Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Desc: iption
_	53	100.0	107	ωį	AAY90824	Aay90824 520C9 hyb
۵	53	00.	243	N	æ	0 52009
ω		100.0	243	N	AAW53170	Aaw53170 520C9 ant
4	53	100.0	243	N	AAW80424	4 520C9
ហ		100.0	243	7	ABW00716	Abw00716 520C9 sFv
6		100.0	246	N	AAR39569	Aar39569 Sequence
7	53	100.0	267	4	AAU04944	Aau04944 Humanised
8	53	100.0	409	4	AAU04945	Aau04945 Humanised
9	53	100.0	534	N	AAR39571	Aar39571 Sequence
10	48	90.6	109	N	AAY06388	Aay06388 Humanised
11	45	84.9	112	S	AAM50879	Aam50879 Anti-bovi
12	43	81.1	11	8	ADM78135	Adm78135 Human SJB
13	43	81.1	90	N	AAR80078	Aar80078 Mouse der
14	43	81.1	92	N	AAW95480	Mouse
15	43	81.1	92	N	AAW95476	
16	43	81.1	107	N	AAR78972	
17	43	81.1	107	N	AAR79247	Aar79247 Light cha
18	43	81.1	107	N	AAR79250	Aar79250 Light cha
19	43	81.1	107	N	AAW27121	Aaw27121 Murine an
20	43	81.1	108	N	AAR21310	Aar21310 Light cha
21	43	81.1	108	8	ADI26702	Adi26702 Mouse ant
22	43	81.1	109	N	AAW18271	7
23	43	•	109	N	AAW85910	0
24	43	81.1	109	4	AAG65862	Aag65862 Anti-PrP
25	43	81.1	109	ហ	ABP51796	Abp5i796 Anti-prio

45	44	43	42	41	40	39	38	37	36	35	34	u u	32	31	30	29	28	27	26
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73.6	73.6	73.6	73.6	73.6	77.4	79.2	79.2	79.2	79.2	79.2	79.2	81.1	81.1	81.1	81.1	81.1	81.1	81.1	81.1
95	95	11	11	11	106	109	108	108	108	108	11	499	257	252	153	144	130	109	109
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AAW85908	AAW18269	ADO60477	AAY44589	AAR52526	ABG76546	AAY06382	AAW23436	AAW23442	AAW23440	AAW23439	AAW23433	AAU72872	AAU72869	AAU72863	ADJ57089	ADJ57086	ADI26651	ADJ10235	ABU58877
Aaw85908	Aaw18269	Ado60477	Aay44589	Aar52526	Abg76546	Aay06382	Aaw23436	Aaw23442	Aaw23440	Aaw23439	Aaw23433	Aau72872	Aau72869	Aau72863	Adj57089	Adj57086	Adi26651	Adj10235	
PrP 81 li	PrP 81 li	Mouse ant	Mouse ant	3D6 light		Humanised	rWI2 ligh	hwi2 ligh	Modified	Modified	CDR-1 of	3B10xP4-1	P5-11 sin		3G4-2BVL-	3G4 antib	20	Murine an	Anti-prio

ALIGNMENTS

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US6054561-A.
                                                                                                                                                                                                              Antigen binding site; immunoglobulin; cancer antigen; immunological; antibody; tumour; human; mucin; cancer; cytostatic; hybridoma; specific binding assay; affinity purification; drug targeting; toxin targeting; imaging; genetic; therapeutic.
                                                                                                                                                                                                                                            520C9 hybridoma VL domain SEQ ID NO:26.
                                                                                                                                                                                                                                                            29-AUG-2000
                                                                                                                                                                                                                                                                        AAY90824;
                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                   AAY90824 standard; protein; 107
                                                                                                                                                                                                                                                           (first entry)
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25-APR-2000.

07-JUN-1995; 95US-00483749

21-MAR-1986; 08-MAY-1988; 11-AUG-1994; 08-FEB-1984; 11-JAN-1985; 84US-00577976. 85US-00690750. 86US-00842476. 88US-00190778.

(CHIR) CHIRON CORP.

94US-00288981

Ring DB;

WPI; 200 N-PSDB;

2000-338508/29. DB; AAA38908.

Monoclonal antibody capable of binding useful for affinity purification, drug treating cancer. 9 6 human breast cancer antigen toxin targeting, imaging, and

Disclosure; Fig 13; 57pp; English.

The present invention describes a monoclonal antibody (MAb) (I) that binds to a human breast cancer antigen that is also bound by MAb 454Cl1 and 520C9 (produced hybridoma ATCC HB8484 and HB8696, respectively). Also described is a hybridoma that produces (I). (I) is useful in specific binding assays, affinity purification, drug or toxin targeting, imaging, and genetic or immunological therapeutics for various cancers. The

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                                                     Variable heavy (VH) and variable light (VL) genes were cloned from a 520C9 hybridoma cDNA library, using probes directed toward the antibody constructed by connecting regions. A two single chain Fv (sFv) gene was constructed by connecting the VH and VL genes with a Ser rich polypeptide linker. The resulting 520C9 two sFv gene, which encodes the present sequence, was inserted into an expression vector, transformed into E. coli, and protein expression induced by the addn. of IPTG to the culture medium. A compsn. comprising a carrier and the 2 sFv protein prod. can be used for in vivo imaging, and drug targetting experiments. The 2 sFv protein prod. is a homodimer, in which both fragments target the same antigen, therefore giving greater binding avidity and longer tissue retention times, compared to individual sFv protein prod. fragments. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Compsns. contg. antigen-targetting antibody fragment constructs comprising dimer of single-chain {\tt Fv} fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence represents a VL domain derived from a 520C9 hybridoma, which is used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Col 33-36; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        520C9; anti-c-erbB-2 monoclonal antibody; single chain Fv; sFv; construct; polypeptide linker; C-terminal amino acid sequence; in vivo imaging; drug targetting experiment; homodimer; increas binding avidity; tissue retention time.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
29-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-333194/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               520C9 anti-c-erbB-2 two single chain Fv construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW02280 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ring DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-00831967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Huston JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Sequence 243 AA,

RESULT 4

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RESULT 3
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XX OF-C
PR 05-C
PR 07-C
PR 07-C
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                                                                                                   Query Match
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Matches 11
                                                                                                                                                                                                                                                                                           This represents the protein sequence of a 520C9 sFv' (single chain Fv) construct. This was constructed by connecting the Vh and Vl genes with a DNA sequence encoding a Serine rich linker. 520C9 is a monoclonal antibody useful in targeting c-erbB-2 antigen. This dimeric construct can be used in the methods of invention of imaging a preselected antigen expressed in a mammal. The methods are used in magnetic resonance imaging of c-erbB-2 or related antigens in cancer diagnosis. The biosynthetic constructs have enhanced properties as in vivo targetting agents in comparison with intact monoclonal antibodies or their Fab fragments. The dimeric constructs permit the in vivo targetting of an epitope on an antigen with greater apparent avidity, including greater tumour specificity, tumour localisation and tumour retention properties than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antigen imaging; single chain Fv; sFv; linker; dimeric; cancer; c-erbB-2; tumour; diagnosis; ss.
                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                       specificity, tumour localisation and tumour retention properti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Col 33-36; 30pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oppermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-FEB-1992;
07-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               520C9 anti-c-erbB-2 sFv'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW53170;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Imaging of antigens in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHIR ) CHIRON CORP.
157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   157 RASQDIGNSLT 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11;
                                                                                                        11;
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RASQDIGNSLT 167
                                                     RASQDIGNSLT 11
                                                                                                                                                                                                                 243 AA;
                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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93US-00133804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-00461838
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                                                                                                      Score 53; DB 2;
Pred. No. 0.057;
Mismatches 0
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Pred. No. 0.057;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of single-chain antibody
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                                                                                                                                                            Length 243;
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AAW80424

ID
AAW80424 standard; pi
XX

AC
AAW80424;

XX

DT
Z8-JAN-1999 (first 6
XX

DT
S20C9 sFv sequence.

XX

S20C9 sFv; antigen; 1
XX

S20C9 sFv; antigen; 1
XX

Synthetic.

XX

Synthetic.

XX

OS
Synthetic.

XX

OS
Synthetic.

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OS
Synthetic.

XX

OS
Synthetic.

XX

CREA-1995; 95US-(
XX

OF
PF
O7-OCT-1993; 93US-(
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CHIR) CHIRON CORP.

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Oppermann H, Houston
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PA
(CHIR) CHIRON CORP.

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Oppermann H, Houston
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PA
(CHIR) CHIRON CORP.

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Oppermann H, Houston
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PT
N-PSDB; AAV63399.

XX

Nucleic acid encodin
PT
and having C-termina
PT
by COCT-1993; 95US-(
AX

CC The present sequence
CC Linker, to produce to complete the invector
CC treatment or diagnos
SQ
Sequence 243 AA;
RESULT 5
ABW00716
ID ABW0
XX ABW0
XX ABW0
XX ABW0
XX BB 5200
XX CEll
KW Cell
KW GFV
XX Unic
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents an antibody 52009 sFV. Variable heavy and light sequences of antibody 752009 are connected, together with a serine linker, to produce the present single chain FV gene. The present sequence exemplifies the invention. Dimers of the single chain FV are used for targeted delivery of drugs or imaging agents (e.g. cytotoxins, prodrugs or 99m-technetium) to antigen-expressing cells, particularly for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid encoding single-chain Fv fragment specific for antigens - and having C-terminal tail for crosslinking to form dimer with improved and having C-terminal tail for crosslinking to form dimer with improved and having contents, used to deliver drugs and imaging agents,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         especially to tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-023541/02.
N-PSDB; AAV63399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            520C9 sPv; antigen; tumour cell; antibody 520C9; targeted delivery; antigen-expressing cell.
    US2002168375-A1
                                                                                 Unidentified
                                                                                                                                                                                                                                                                         520C9 sFv protein.
                                                                                                                                                                                                                                                                                                                                                           15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                    ABW00716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABW00716 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Col 35-36; 29pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CREA-) CREATIVE BIOMOLECULES INC (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                               proliferation; cytotoxic agent; drug targetting; single-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 RASQDIGNSLT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RASQDIGNSLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; |
|llarity 100.0%; |
|Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosis
                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-00831967.
93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-00461386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ľ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 53; DB 2;
Pred. No. 0.057;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (especially of ovary or breast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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RESULT 6
AAR39509
ID AAR397
XX AAR397
AC AAR3
XX AAR397
DT 25-N
DT 07-F
XX Sequ
XX Sing
XX Sing
XX Sing
XX W093
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a binding protein formulation for targetting an epitope on an antigen expressed in mammal. The formulation comprises of dimeric blosynchetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain. The invention is useful for imaging a preselected antigen in a mammal expressing the antigen. It is also useful for targetting drugs that inhibits cell proliferation and cyrotoxic agents that kill cells. The present sequence is 52009 sFv protein. This sequence is used in the exemplification of the
                                                                    (CREA-)
                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                      Single chain
biosynthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
07-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel binding protein formulation for targeting epitope on antigen in mammal, comprising dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-OCT-1993; 93US-00133804
05-JUN-1995; 95US-00462641
26-APR-2000; 2000US-00558741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-NOV-2002
                     Huston JS,
                                                                                                                                        06-FEB-1992;
                                                                                                                                                                                    05-FEB-1993;
                                                                                                                                                                                                                                     19-AUG-1993
                                                                                                                                                                                                                                                                                 WO9316185-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence of 520C9 sFv protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR39569;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR39569 standard; protein; 246 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 19-20; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Huston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2003-765156/72.
)B; AAD61485.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 RASODIGNSLT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
                                                                    CETUS ONCOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RASQDIGNSLT 167
                     Houston LL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Houston
                                                                                                                                                                                                                                                                                                                                                                         Fv polypeptide; VH-VL heterodimer; immunoglobulin; single polypeptide chain binding site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-00887853.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                        92US-00831967.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92US-00831967.
93US-00133804.
                                                                                                                                                                                    93WO-US001055
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                     Ring
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                            INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 53; DB 7;
Pred. No. 0.057;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB,
                     DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                     Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>.</u>
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RESULT 7
AAU04944
ID AAU04949
ID AAU04949
XX AAU0
AC AAUC
XX AAUC
AX CAAUC
XX CAAUC
COS Homa
XX CAAUC
COS Homa
XX CAAUC
COS Homa
XX CAAUC
COS CAAUC
FT Regi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                                                                                                  Mus sp.
Synthetic.
Chimeric.
                                                                                                                      Region
                                         Region
                                                                                Region
                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cerbB-2 refers to a protein antigen expressed on the surface of tumour cells. such as breast and ovarian tumour cells, which is an approx.

200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). A single chain Fv (sFv) is a covalently linked VH-VL heterodimer which is expressed from a gene fusion including VH- and VL- encoding genes connected by a peptide-encoding linker. Such linker sequences are set forth in AA residues 116-135 in AAR39569, which includes part of the 16 AA linker sequences in AAR39572. Using AAQ46084 for the 520C9 monoclonal antibody, a single chain polypeptide can be produced having a binding affinity for a c-erbB-2 related antigen. 'X' in AAR39569 refers to the location of a stop codon in AAQ46084. (Updated on 55-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                               Humanised; antibody; p185; 520C9H; interleukin-2; IL-2; immunoconjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Humanised anti-p185 single chain antibody, 520C9H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                       cancer; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU04944;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New single chain Fv imaging or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU04944 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1993-272889/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RASQDIGNSLT 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RASQDIGNSLT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA,
                                                                                                                                                                                                                                                                                                                                                                                                        adenocarcinoma
                                                 note= "Complementarity determining
                                                                                                                                                                                                                  /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                         'label= CDR
'note= "Complementarity determining
                                                                                                                                                                                                                                                          /label= CDR /note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                    label= CDR
note= "Complementarity determining
                                                                                                                                     note= "Links
                                                                                                                                                  88. .152
|abel= Synthetic
                                                                   abel= CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                       .249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptide binding to C-erbB-2 tumour antigen - breast or ovarian cancer etc.
                                                                               .208
                                                                                                                                                                                                          .126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                         .186
       "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 53; DB Pred. No. 0.0; Mismatches
                                                                                                                                      the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A
                                                                                                                                  _peptide_linker
heavy chain to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2; Le. 0.057;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 246;
                                                                                                                                   the
                                                                                                                                                                           region'
                                                                                            region"
                                                                                                                                  light chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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ACC CAN ACC CA
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AAU04945
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Region
                                           Protein
                                                                                                                                                             Key
                                                                                                                                                                                                                              Mus sp.
Synthetic.
                                                                                            Protein
                                                                                                                                       Peptide
                                                                                                                                                                                                         Chimeric.
                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                        cancer;
                                                                                                                                                                                                                                                                                                                                              Humanised;
                                                                                                                                                                                                                                                                                                                                                                                       Humanised anti-p185 antibody/IL-2 fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                        06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU04945;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU04945 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence represents a humanised anti-p185 single chain antibody which is linked to a human interleukin-2 (IL-2) molecule to make a fusion protesin. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cancers particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, renal system, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel immunoconjugate comprises a humanized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 267 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gastrointestinal tract or gastric tumours.
correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 7; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Austin R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JAN-2001; 2001WO-US001919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JUL-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHIR ) CHIRON CORP.
(HAMI-) HAMILTON CIVIC HOS
(HAMI-) HAMILTON REGIONAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JAN-2000; 2000US-0177258P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 RASODIGNSLT 11
                                                                                                                                                                                                                                                                                                                        tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS09507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RASODIGNSLT
                                                                                                                                                                                                                                                                                                                                                antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9; 74pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
/label=
53. .57
                                                          23. .409
/label=_Mature_fusion_protein
                                                                                                                                                                                                                                                                                                                      pody; p185; 520C9H; interleukin-2;
adenocarcinoma; fusion protein.
                                                                                                        /label= Signal_peptide
                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              260. .267
/label= Glu_Glu_epitope
                                              . 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful for inhibiting tumor cell growth in vivo anti-p185 antibody linked to an Interleukin-2
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                Humanised_antibody_520C9H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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0.063;
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                                                                                                                                                                                                                                                                                                                                            IL-2; immunoconjugate;
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Best Local S
Matches 11
                                                                                                           The sequence represents a humanised anti-p185 (520C9) single chain antibody/human interleukin-2 (IL-2) fusion protein. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumencurs or cances particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, rensestem, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                         polypeptide.
                                                                                                                                                                                                                                                                       WPI; 2001-451904/48.
N-PSDB; AAS09508.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                     Novel immunoconjugate comprises a humanized
                                                                                                                                                                                                                                                                                                         Austin R, Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                                                   19-JAN-2001; 2001WO-US001919
                                                                                       Sequence
                                                                                                                                                                                                    Claim 13; Fig 11; 74pp; English
                                                                                                                                                                                                                                                                                                                             (CHIR ) CHIRON CORP.
(HAMI-) HAMILTON CIVIC HOSPITALS
(HAMI-) HAMILTON REGIONAL CANCER
                                                                                                                                                                                                                                                                                                                                                                            20-JAN-2000; 2000US-0177258P
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176
                     ب
                                          Similarity 100
11; Conservative
                                                                                       409
 RASQDIGNSLT
                    RASODIGNSLT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /label= Synthetic_linker peptide
/note= "Links the antibody to the
277. .409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'note= "Encode by ACA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'label= CDR
'note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         label IL_2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               label CDR note "Complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             label= CDR note= "Complementarity determining region"
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note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    label= Synthetic_peptide_linker
note= "Links the heavy chain to the light chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "Encoded by GGG"
                                                     100.0%;
186
                                                                                                                                                                                                                                                                                                         Ring DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .249
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                                                                                                                                                                                                                                     useful for inhibiting tumor cell growth in vivo anti-pl85 antibody linked to an Interleukin-2
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                                            0
                                                     Score 53; DB Pred. No. 0.1;
                                            Mismatches
                                                                                                                                                                                                                                                                                                                               CENT.
                                                                                                                                                                                                                                                                                                                                          RES DEV INC
                                                               DB 4;
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                                                                 Length 409;
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RESULT 9

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RESULT 10
AAY06388
ID AAY06388
AC AAY06
XX AAY06
XX AAY06
XX Human
XX Human
XX Monooc
KW Monooc

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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                      Key
Peptide
                                                                                                                                                                                  Humanised antibody; antibody humanisation; antibody engineering; LM609; monoclonal antibody; complementarity determining region; CDR grafting; mouse; human; integrin; apoptosis; angiogenesis; cancer; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. such as breast and ovarian tumour cells, which is an approx. 200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New single chain Fv polypeptide binding to C-erbB-2 tumour antigen - imaging or treating breast or ovarian cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-FEB-1992;
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07-FEB-1994
                                                                                           Synthetic
                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                   Humanised LM609 antibody VL
                                                                                                                                                                                                                                                                                                                                  06-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                  AAY06388;
                                                                                                                                                                                                                                                                                                                                                                                                                              AAY06388 standard; protein; 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 534
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAQ46086.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1993-272889/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Huston JS, Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-FEB-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CREA-) CREATIVE BIOMOLECULES (CETU) CETUS ONCOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 448 RASQDIGNSLT 458
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                   Location/Qualifiers
1...2
/note= "vector-encoded residues"
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Pred. No. 0.13;
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RESULT 11
AAM50879
ID AAM50
XX AAM50
XC AAM50
AC AAM50
DT 07-MA
XX Prion
KW Infec
XX Infec
XX Wus n
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XX W020(
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                             27-JUL-2000; 2000US-00627218.
                                                                                07-FEB-2002.
                                                                                                        WO200210335-A2
                                                                                                                                 Mus musculus
                                                                                                                                                     Prion; prion protein; PrP; PrPC; PrPSc; antibody; Fab; mouse; ungulate; infection; bovine spongiform encephalopathy; BSE; diagnosis; therapy.
                                                                                                                                                                                                                       07-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents the light chain variable region of a humanised LM609 antibody. LM609 is directed to human integrin alpha-v beta-3. It selectively promotes apoptosis of vascular cells that have been stimulated to undergo angiogenesis, making it a tool for cancer diagnosis and therapy. The invention provides humanised antibodies, especially humanised LM609. In such humanized antibodies, a light chain, CDR from a mouse antibody such as LM609 is grafted onto a human light chain, and a heavy chain CDR from a mouse antibody is grafted onto a human antibody heaving the desired specificity is selected. By preserving the original CDR sequences such as the HCDR3 and LCDR3 sequences of LM609 (see
      (REGC ) UNIV CALIFORNIA
                                                     17-JUL-2001; 2001WO-US022648.
                                                                                                                                                                                            Anti-bovine prion protein peptide antibody light chain.
                                                                                                                                                                                                                                                AAM50879;
                                                                                                                                                                                                                                                                        AAM50879 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 109 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 52; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Production of humanized mouse monoclonal antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-394979/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                       24
                                                                                                                                                                                                                                                                                                                                                                                         10;
                                                                                                                                                                                                                                                                                                                                                               1 RASQDIGNSL 10
                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                       RASODIGNSL
                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rader C;
                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-00986016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-US025828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "CDR1"
50. .56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "CDR3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       50. .56
/note= "CDR2"
                                                                                                                                                                                                                                                                       protein; 112
                                                                                                                                                                                                                                                                                                                                       ű
                                                                                                                                                                                                                                                                                                                                                                                                  90.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . 97
                                                                                                                                                                                                                                                                                                                                                                                      Score 48; DB;
; Pred. No. 0.2:
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                        B
                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                              Length 109
                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                      0
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Sequence 112 AA;

as

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CC recombinant attibody fragments (Fabs) were isolated from mouse cDNA and Cc cloned into a vector that expressed human-mouse chimeric Fabs in CC Escherichia coli. The purified Fabs were labeled with Europium, giving an CC Escherichia coli. The purified Fabs were labeled with Europium, giving an CC Fab designated Eu-(HuW) Fab P. This is an example of antibodies of the CC invention that specifically bind with a high degree of binding affinity CC to a native ungulate PrPC in situ and/or a denatured ungulate PrPSC, but to a native ungulate PrPSC in situ. The antibodies may specifically bind to epicopes of PrPC of a specific species of animal or to 1 or more types of PrPC proteins from 1 or more species of animal or to 1 or more CC types of PrPC proteins from 1 or more species of ungulates. The antibodies are useful for the detection of prion infection in ungulates, particularly cattle. A fast, efficient, cost-effective assay is achieved. The assay can be used to screen for the presence of PrPSC in products such as pharmaceuticals (derived from natural sources), food, cosmetics etc. The antibodies can be used with a compound that denatures PrPSC thereby providing a means of differentiating levels of PrPC and PrPSC in a sample. The invention also provides a therapeutic antibody which prevents or treats prion disease in ungulates, and correctifically in Cartia and a means of for certificing to ungulates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ELISA. The presence of PrPSc in tissues of humans or animals is indicative of prion infection. The Fab was generated against the 96-105 region of bovine prion protein (PrP). Clone P binds to an epitope (see AAM50886-87) that is present in all ungulate PrP sequences, including bovine, and also in other species such as human, sheep and mouse. The P
                                                  being prion free
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel antibody useful for detecting infectious prions, particularly pathogenic scrapie isoform of the prion protein, PrPSc in an ungulate, especially cow, binds to native ungulate prion protein C in situ.
                                                                                   specifically in cattle, and a means
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is that antibody fragment (Clone P) t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 41; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prusiner SB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SCRI ) SCRIPPS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sent sequence is that of the light chain of a recombinant fragment (Clone P) that binds tightly to denatured boving to the native conformation of the same protein in CDIO for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Safar J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Williamson AR,
                                                                          for certifying certain products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Burton DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        formatted
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Matches
                                                                                                                                                                                                                                                        Query Match
                                                                         biochemical;
                       08-APR-2004.
                                       WO2004029094-A1
                                                                         complementarity determining r
biochemical; human; variable;
                                                                                         monoclonal antibody; adenylate kinase isozyme
                                                                                                          Human SJB3-86 antibody variable light chain CDR1 peptide region.
                                                                                                                            01-JUL-2004
                                                                                                                                            ADM78135;
                                                                                                                                                           ADM78135 standard; peptide; 11
                                                                                                                                                                                                                                                Local
                                                        sapiens.
                                                                                                                                                                                                       24
                                                                                                                                                                                                                      1 RASODIGNSL 10
                                                                                                                                                                                                                                       Similarity 9; Conserv
                                                                                                                                                                                                       RASQDIGNNL
                                                                                                                                                                                                                                        Conservative
                                                                         human;
                                                                                                                          (first entry)
                                                                                                                                                                                                                                              84.9%;
                                                                                                                                                                                                                                       Score 45; DB 5
Pred. No. 0.9;
1; Mismatches
                                                                                  region;
                                                                                                                                                                                                                                       1;
                                                                        gion; CDR; heart disease; marker AK3
light chain.
                                                                                                                                                            A
                                                                                                                                                                                                                                                       5
                                                                                         u
                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                       Length 112
                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                      Gaps
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27-SEP-2003; 2003WO-KR001979.

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ARESULT 13
AARROD AARROD
XX AARROD
XX AARROD
XX AARROD
XX MOUSE
XX Light
XX Light
XX Key
FT Regic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel monoclonal antibody specific to adenylate kinase isozyme 3, comprising 4 or more of 6 complementarity determining regions (CDRs) of a sequence having 6-18 amino acids. The invention further relates to: a composition comprising the monoclonal antibody for detecting a heart disease marker AK3; a kit comprising the monoclonal antibody for the diagnosis of heart disease; and a method of detecting a heart disease marker AK3. The monoclonal antibody is useful for preparing a composition for detecting a heart disease marker AK3. The monoclonal antibody reduces false positive results of the conventional biochemical markers. This sequence represents a human antibody variable light chain CDR peptide region of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New monoclonal antibody specific to human mitochondrial adenylate kinase isozyme 3, useful for preparing a composition for detecting a heart disease marker \lambda K3 .
               30-MAR-1994;
                                                                                                                                                                                                             Regior
                                                                                                                                                                                                                                           Region
                                                                                                                                                                                                                                                                                                                                                   Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                              Light chain; RT3; murine; catalytic antibody; bacteriophage; pattern
                                                                                                                                                                                                                                                                                                                                                                                                          Mouse derived light chain RT3 phage antibody pattern A.
                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAY-1996 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR80078;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR80078 standard; protein; 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
                                             12-OCT-1995.
                                                                                                                      Region
                                                                                                                                                  Region
                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                       Region
                                                                                                                                                                                                                                                                                                     Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-316088/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-SEP-2002; 2002KR-00059211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KIMH/) KIM H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   μ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 90.
9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RASQDIGNSL 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 98; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RASQDIGSSL 10
               94WO-US003420
                                                                                                        89. .90
/note=
                                                                                                                                    note-
                                                                                                                                                                                              note=
                                                                                                                                                                                                                           /note= "framework region 2"
                                                                                                                                                                                                                                                         note
                                                                                                                                                                                                                                                                                     'note= "framework region 1"
                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                . 47
                                                                                                                                                                                  . 78
                                                                                                                                                                                                                                                                         . 26
                                                                                                                                                                                                                                            .40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        81.1%;
90.0%;
                                                                                                                                                    . 88
                                                                                                                                                               "framework region 3"
                                                                                                                                                                                             "complementarity determining
                                                                                                                                                                                                                                                       "complementarity determining
                                                                                                      "framework region 4 N-terminal fragment"
                                                                                                                                  "complementarity determining region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 43; DB 8;
Pred. No. 0.18;
L; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₿
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                             region
                                                                                                                                                                                                                                                       region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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Best Local Similarity
Matches 9; Conserv
                                                                                 (MCCA/) MCCAFFERTY J.
(CHIS/) CHISWELL D.
(DARS/) DARSLEY M J.
(TITW/) TITWAS R C.
(MART/) MARTIN M T.
(MART/) MARTIN J H.
(KENT/) KENTEN J H.
(SMIT/) SMITH R.
(FITZ/) FITZGERALD K.
(WILL/) WILLIAMS R O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT04625 encodes AAR80078 mouse derived light chain RT3 phage antibody. The DNA was used in the prepn. of catalytic antibody (CA) producing bacteriophage. The CAs can be used to activate/deactivate a biological function in an animal by enhancing the rate of cleavage, or formation of a specific bond within a mol. in vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Smith RG,
Kenten JH,
                                                                                                                                                                                                                                                                                                Catalytic; antibody; phage display; immunising; phage expression vector; prodrug; scFV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Production of catalytic antibodies displayed on phage - gene library of antibody-derived domains and expressing
                     N-PSDB; AAX00879.
                                WPI; 1999-105036/09
                                                    Fitzgerald K, Darsley
Kenten JH, Chiswell D,
                                                                                                                                                                                                              14-JUL-1994;
                                                                                                                                                                                                                                   05-JAN-1999.
                                                                                                                                                                                                                                                        US5855885-A.
                                                                                                                                                                                                                                                                           Mus sp
                                                                                                                                                                                                                                                                                                                                  Mouse derived RT3 phage antibody light chain pattern C genetic sequence
                                                                                                                                                                                                                                                                                                                                                       29-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                            AAW95480;
                                                                                                                                                                                                                                                                                                                                                                                                AAW95480 standard; protein; 92 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 90 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 9; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-358624/46.
N-PSDB; AAT04625.
                                                                                                                                                                                          22-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IGEN-) IGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RASQDIGNSL 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                     RASODIGSSL 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mccafferty J, Chiswell D, Darsley MJ, Martin MT, Titmas RC, Williams RO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                          93US-00007684
                                                                                                                                                                                                               94US-00273146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94WO-US003420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      81.1%;
                                            MJ, Williams, Mccafferty J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 43; DB Pred. No. 1.8; Nismatches
                                                              Williams RO,
                                                     Titmas RC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fitzgerald
                                                               Martin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          it by
                                                                3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         generating a 
in phage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   χ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Production of catalytic antibodies displayed on bacteriophages

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cc displayed on a phage. The method comprises: (a) generating a gene library cf antibody-derived domains; (b) inserting coding for the domains into a cc phage expression vector; and (c) isolating the catalytic antibodies. The cx phage expression vector incorporates a histidine peptide in tandem with a cc myc peptide. The catalytic antibodies can be isolated by preparing an cc antigen; optionally immunising an animal with the antigen; generating a cc library of VH and VL domains from the immunised animal; cloning the VH cr and VL domains into a phage expression vector to generate phage display cc antibodies; selecting phage display antibodies which bind specifically to the antigen; screening the selected phage display antibodies for catalytic activity to substrate; and isolating the catalytic antibodies, which can be used for in vivo activation of a prodrug. The cr tandem with a myc peptide. The processes are used to produce catalytic antibodies, which can be used for in vivo activation of a prodrug. The present sequence represents a genetic sequence of light chain PCR pattern cc from mouse derived RT3 phage antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 15
AAW95476
Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Fig 11; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprises generating coding into a phage e
                                          Fitzgerald K, Darsar
TH, Chiswell
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Catalytic; antibody; phage display; immunising; phage expression vector; prodrug; scPV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 92
                                                                                                                                                                                          (CHIS/)
(DARS/)
(TITM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse derived RT3 phage antibody light chain pattern A genetic sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW95476;
   N-PSDB; AAX00875.
                                                                                                                                                                                                                                                                                                                                                      05-JAN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                         ds snw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW95476 standard; protein; 92 AA.
                    WPI; 1999-105036/09
                                                                                                                                                                                                                                                                                    22-JAN-1993;
                                                                                                                                                                                                                                                                                                                     14-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                       US5855885-A.
                                                                                                                                                           (KENT/
                                                                                                                                                                                                                                               (MCCA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16 RASQDIGSSL
                                                                                                        SMITH R.
FITZGERALD K.
WILLIAMS R O.
                                                                                                                                                       CHISWELL D.
DARSLEY M J.
TITMAS R C.
MARTIN M T.
KENTEN J H.
                                                                                                                                                                                                                                               MCCAFFERTY J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RASQDIGNSL 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ĀĀ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                     Darsley MJ, Williams RO, Smith R, iswell D, Mccafferty J, Titmas RC;
                                                                                                                                                                                                                                                                                                                     94US-00273146.
                                                                                                                                                                                                                                                                                    93US-00007684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a gene library of antibody-derived domains inserting expression vector and isolating the catalytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 43; DB Pred. No. 1.8; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for producing catalytic antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                           Martin MT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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Production of catalytic antibodies displayed on bacteriophages comprises generating a gene library of antibody-derived domains inserting coding into a phage expression vector and isolating the catalytic
                                                                                                                                                    antibodies.
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Example 4; Fig 9A-F; 117pp; English.

The invention relates to methods for producing catalytic antibodies CC displayed on a phage. The method comprises: (a) generating a gene library CC of antibody-derived domains; (b) inserting coding for the domains into a CC phage expression vector; and (c) isolating the catalytic antibodies. The CC phage expression vector incorporates a histidine peptide in tandem with a CC myc peptide. The catalytic antibodies can be isolated by preparing an CC antigen; optionally immunising an animal with the antigen; generating a CC library of VH and VL domains from the immunised animal; cloning the VH CC and VL domains into a phage expression vector to generate phage display antibodies which bind specifically to CC the antigen; screening the selected phage display antibodies for CC thatlytic activity to substrate; and isolating the catalytic antibodies, which a myc peptide. The processes are used to produce catalytic cantibodies, which can be used for in vivo activation of a prodrug. The CC present sequence represents a genetic sequence of light chain pattern A CC from mouse derived RT3 phage antibodies

Sequence 92 AA;

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밁
            Š
                            Query Match
Best Local S
Matches 9
16
              1 RASODIGNSL 10
                             Similarity 9; Conserv
RASODIGSSL
                             Conservative
25
                                     81.1%;
                              Score 43; DB Pred. No. 1.8; 1; Mismatches
                                             ۲.
                               0,
                                             Length 92;
                                 Indels
                                0
                               Gaps
                                 0
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Search completed: April Job time : 28.9649 secs

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2005, 15:47:15

Sequence 4, Appli Sequence 67, Appli Sequence 67, Appli Sequence 73, Appli Sequence 22, Appli Sequence 22, Appli Sequence 24, Appli Sequence 28, Appli Sequence 28, Appli Sequence 29, Appli Sequence 21, Appli Sequence 22, Appli Sequence 23, Appli Sequence 23, Appli Sequence 24, Appli Sequence 28, Appli Sequence 28, Appli Sequence 30, Appli Sequence 30, Appli Sequence 30, Appli Sequence 30, Appli Sequence 31, Appli

Sequence 3, Appli Sequence 72, Appl Sequence 72, App

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Result
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein - protein search, using
 111098765432
                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                       Score
                                                                                                                                                                                                                                                                                                                            Published Applications AA:*

1: //ggn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: //ggn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

3: //ggn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

4: //ggn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

5: //ggn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

6: //ggn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

7: //ggn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

8: //ggn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

9: //ggn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

10: //ggn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

11: //ggn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

12: //ggn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

13: //ggn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

14: //ggn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

15: //ggn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

16: //ggn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

16: //ggn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*

18: //ggn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep:*

19: //ggn2_6/ptodata/1/pubpaa/US10B_PUB.pep:*

19: //ggn2_6/ptodata/1/pubpaa/US10B_PUB.pep:*
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length: 2000000000
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Match
            100.0
100.0
100.0
100.0
90.6
84.9
81.1
81.1
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Gapop 10.0 ,
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                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd
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          7 US-10-683-547-6

US-09-766-543-1.0

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4 US-10-078-7578-55

6 US-10-803-622-267

6 US-10-803-622-267

6 US-10-803-623-267

6 US-10-943-965-3-267

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       Sequence 6, Appli
Sequence 6, Appli
Sequence 10, Appl
Sequence 12, Appl
Sequence 15, Appl
Sequence 45, Appl
Sequence 267, App
Sequence 267, App
Sequence 74, Appl
Sequence 74, Appl
Sequence 74, Appl
Sequence 74, Appl
Sequence 75, Appl
                                                                                                                                                                                                     Description
                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1
US-09-887-853-6
Sequence 6, Application US/09887853; Sequence 10. US20020168375A1
PALENT NO. US20020168375A1
GENERAL INFORMATION:
James S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

REDIUM TYPE: Floppy disk

COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHTIN Release #1.0, Ver
CURRENT APPLICATION UNMBER: US/09/887,853
FILING DATE: 21-Jun-2001
CLASSIFICATION UNMBER: US/09/887,853
FILING DATE: 21-Jun-2001
CLASSIFICATION UNMBER: US/08/133,804
FILING DATE: CURKnown>
PRIOR APPLICATION NUMBER: US/08/133,804
FILING DATE: CURKnown>
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION UNMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwit
STREET: Exchange Place,
                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     73336622222
                                                                                                                                                                                                                                                                                                                                                                                            Oppermann, Hermann
Houston, L. L.
Ring, David B.
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6. US-10-620-850-4
6. US-10-642-116-4
7. US-10-642-1058-4
7. US-10-642-128-4
7. US-10-642-128-4
7. US-10-239-656-67
5. US-10-239-656-67
5. US-10-239-656-73
7. US-10-808-538-4
7. US-10-808-538-4
7. US-10-808-538-23
7. US-09-155-106-24
7. US-09-155-106-28
7. US-09-155-106-28
7. US-09-155-106-28
7. US-09-155-106-28
7. US-09-155-106-38
7. US-10-808-538-23
7. US-10-808-538-23
7. US-10-808-538-23
7. US-10-808-538-38
7. US-10-808-538-38
7. US-10-808-538-38
7. US-10-808-538-38
7. US-10-808-538-38
8. US-10-778-38
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Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-642-120-4
US-10-642-060-4
US-10-642-122-4
US-10-642-124-4
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TELEPHONE: 617-248-7477

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US-09-766-543-10
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NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                         Sequence 10, Application US/09766543
Patent No. US20020041865A1
                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                   APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: King, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PP01679.002
CURRENT APPLICATION NUMBER: US/09/766,543
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Oppermann, H.
TITLE OF INVENTION: BOYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING
FILE REFERENCE: CIBT-PO1-130
CURRENT APPLICATION NUMBER: US/10/683,547
CURRENT FILING DATE: 2003-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US/09/558,741
PRIOR FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 07/831,967
PRIOR FILING DATE: 1992-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Huston, J.
APPLICANT: Houston, J
APPLICANT: Ring, D.
APPLICANT: Oppermann.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-248-7100 INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                      157 RASODIGNSLT 167
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Pred. No. 0.046;
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GENERAL INFORMATION:

APPLICANT: BARBAS, Carlos F., III

APPLICANT: RADER, Christoph

ITITLE OF INVENTION: HUMANIZATION OF MURINE ANTIBODY

FILE REPERENCE: TSRI 598.0 Con1

CURRENT APPLICATION NUMBER: US/10/078,757B

CURRENT FILING DATE: 2002-02-19

PRIOR APPLICATION NUMBER: US 08/986,016

PRIOR FILING DATE: 1997-12-05

NUMBER OF SEQ ID NOS: 56

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 55
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US-10-078-757B-55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Ring, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PP01679.002
                                                                                                                                                                                                                                                                                                                                         Sequence 55, Application US/100 Publication No. US20030166871A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/766,543
CURRENT FILING DATE: 2000-01-19
                      LENGTH: 109
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: H520C9sFv plus OTHER INFORMATION: linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: 520C9 OTHER INFORMATION: humanized single-chain antibody used in the OTHER INFORMATION: IL-2-antibody fusions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: hybrid mouse - human sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 267
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             176 RASQDIGNSLT 186
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100.0%; F
itive 0;
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Pred. No. 0.052;
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Pred. No. 0.051;
; Mismatches
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APPLICANT: Burton, Dennis

TITLE OF INVENTION: Antibodies Specific for Ungulate PrP
FILE REFERENCE: UCAL-194
CURRENT APPLICATION NUMBER: US/10/355,780
CURRENT FILING DATE: 2003-01-30
PRIOR APPLICATION NUMBER: US/09/627,218B
PRIOR FILING DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 112
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthesized peptide
US-10-355-780-1
                           APPLICANT: Miller, David J.
APPLICANT: Miller, David J.
APPLICANT: Miller, David J.
APPLICANT: Miller, David J.
APPLICANT: Pease, Larry R.
APPLICANTON: System
FILLE REFERENCE: 1199-1-005CIP2
CURRENT APPLICATION NUMBER: US/10/010,729
CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 09/33,473
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR FILING DATE: 1000-05-28
PRIOR APPLICATION NUMBER: 09/322,862
PRIOR APPLICATION NUMBER: 09/322,862
PRIOR APPLICATION NUMBER: 08/779,784
PRIOR FILING DATE: 1997-01-07
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR PILING DATE: 1996-08-08
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR PILING DATE: 1996-08-08
PRIOR PILING DATE: 1996-08-08
PRIOR APPLICATION NUMBER: 08/236,520
PRIOR PILING DATE: 1996-04-29
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Publication No. US20030143224A1
GENERAL INFORMATION:
APPLICANT: Prusiner, Stanley
APPLICANT: Safar, Jiri
APPLICANT: Williamson, Anthony
APPLICANT: Burton, Dennis
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Best Local 8
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for Windows Version
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90.0%;
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Pred. No. 0
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OTHER INFORMATION: light chain from clone MIF US-10-803-622-267
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US-10-803-622-267
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; ORGANISM: Mus musculus
US-10-010-729-45
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; LENGTH: 108
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Best Local Similarity 90.0%;
Matches 9; Conservative
                  Query Match
Best Local
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  Matches
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CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
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                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
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PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR FILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: GB 9015198.6 PRIOR FILING DATE: 1990-07-10 PRIOR APPLICATION NUMBER: GB 9022845.3 PRIOR FILING DATE: 1990-10-19
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APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
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                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: GB 9110549.4 PRIOR FILING DATE: 1991-05-15
                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                         ENGTH: 108
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                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/971,857
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l Similarity
9; Conserv
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Cambridge Antibody Technology Limited
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Griffithe, Andrew
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  Conservative
                  81.1%;
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Score 43; DB 16; Length 108; Pred. No. 1.7; 1; Mismatches 0; Indels
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Pred. No. 1.7;
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24 RASQDIGSSL 33

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CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILLING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILLING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PRILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 9110549.4
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                                                                               Sequence 74, Application U
Patent No. US20020150571A1
GENERAL INFORMATION:
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Best Local
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SEQ ID NO 267
LENGTH: 108
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APPLICANT:
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PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1995-06-07
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APPLICANT: Cambridge Antibody Technology Limited
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TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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Local Similarity 90.0%;
Williamson, R. Anthony
Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/971,857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/GB91/01134 FILING DATE: 1991-07-10
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                                                           APPLICANT: Prusiner, Stanley B.
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RASQDIGSSL 33
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Griffiths, Andrew
Jackson, Ronald
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Chiswell, David
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b. US20040157215A1
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PRIOR APPLICATION NUMBER: 09/943,906
PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/550,374
PRIOR PRILING DATE: 2000-04-13
PRIOR APPLICATION NUMBER: 09/036,579
PRIOR APPLICATION NUMBER: 09/036,579
PRIOR APPLICATION NUMBER: 09/713,939
PRIOR FILING DATE: 1996-09-13
PRIOR APPLICATION NUMBER: 08/528,104
PRIOR APPLICATION NUMBER: 08/528,104
PRIOR FILING DATE: 1995-09-14
NUMBER OF SEQ ID NOS: 86
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 74, Application US/104
Publication No. US20030228303A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                APPLICANT: Prusiner, Stanley B.
APPLICANT: Williamson, R. Anthony
APPLICANT: Burton, Dennis R.
TITLE OF INVENTION: Antibodies Specific for Native PrPSc
FILE REFERENCE: UCAL059CON3
                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/435,602
CURRENT FILING DATE: 2003-05-09
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INFORMATION FOR SEQ ID NO: 74:
Q ID NO 74
LENGTH: 109
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 74:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Bozicevic, Karl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24 RASQDIGSSL 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RASODIGNSL 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/943,906 FILING DATE: 30-Aug-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 28,807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/10435602
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90.0%;
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Pred. No. 1.7;
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46 RASQDIGSSL 55

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RESULT 12
US-08-779-784-35
; Sequence 35, Application US/08779784
; Publication No. US20020164325A1
; GENERAL INFORMATION:
ROGRIGUEZ, MOSES
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; ORGANISM: mouse
US-10-435-602-74
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Best Local Similarity
Marches 9; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/779,784
PILING DATE: 07-JAN-1997
CLASSIFICATION: 424
PRICR APPLICATION DATA:
APPLICATION NUMBER: US 08/692,084
PILING DATE: 08-AUG-1996
PRICR APPLICATION NUMBER: US 08/692,084
APPLICATION NUMBER: US 08/692,084
PILING DATE: 08-AUG-1996
PRICR APPLICATION NUMBER: US 08/692,084
APPLICATION NUMBER: US 08/692,084
                                      Matches
                                                                       Query Match
                                                                                                                                                                                                                                                          TELEFAX: 201-343-1684
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 amino acids
TYPE: amino acid
                                                                                                                                                 HYPOTHETICAL: N
FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Rodriguez, Moses
APPLICANT: Miller, David J.
APPLICANT: Asakura, Kunihiko
TITLE OF INVENTION: PROMOTION OF CENTRAL NERVOUS SYSTEM
TITLE OF INVENTION: REMYELINATION USING MONOCLONAL AUTOANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 29-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                            NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1199-1-001 CIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Hackensack
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: David A. Jackson, Esq.
STREET: 411 Hackensack Ave, Continental Plaza, 4th
STREET: Ploor
                                                                                                                                ORGANISM:
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1 RASQDIGNSL 10
                                      Similarity
9; Conserv
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linear
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90.0%;
                                                      81.1%;
                                    Score 43; DB Pred. No. 2; 1; Mismatches
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APPLICANT: Miller, David J.
APPLICANT: Miller, David J.
APPLICANT: Miller, David J.
APPLICANT: Pease, Larry R.
ITILE OF INVENTION: Human IgM Antibodies and Diagnostic and
ITILE OF INVENTION: Therapeutic Uses Thereof Particularly in the Central Nervous
ITILE OF INVENTION: System
FILE REFERRNCE: 1199-1-005CIP2
CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: US/10/010,729
CURRENT FILING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: 09/730,473
PRIOR APPLICATION NUMBER: 09/30,473
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR APPLICATION NUMBER: 09/522,862
PRIOR APPLICATION NUMBER: 08/79,784
PRIOR FILING DATE: 1999-05-28
PRIOR APPLICATION NUMBER: 08/692,084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1994-04-29

NUMBER OF SEQ ID NOS: 80

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 71

LENGTH: 130

TYPE: PRT

ORGANISM: Mus musculus

US-10-010-729-71
                                                                                  ; LENGTH: 144
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-642-120-4
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                                                                                                                                                                                                    SEQ ID NO 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/10642120 Publication No. US20040131610A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
  Best Local Similarity
                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                   APPLICANT: Thorpe, Philip B.
APPLICANT: Thorpe, M. Melina
APPLICANT: Soares, M. Melina
APPLICANT: Ran, Sophia
TITLE OF INVENTION: Methods for Treating Viral Infections Using Antibodies to
TITLE OF INVENTION: Aminophospholipids
FILE REFERENCE: 4001.002900
CURRENT APPLICATION NUMBER: US/10/642,120
CURRENT FILING DATE: 2003-08-15
PRIOR APPLICATION NUMBER: US 10/621,269
PRIOR PILING DATE: 2003-07-15
PRIOR APPLICATION NUMBER: 60/396,263
PRIOR FILING DATE: 2003-07-15
                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.1
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Pred. No. 2
  Score 43;
Pred. No.
     DB 16;
2.3;
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                             Length 144;
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US-10-642-060-4

Sequence 4, Application US/10642060

Publication No. US20040131621A1

GENERAL INFORMATION:

APPLICANT: Thorpe, Philip E.

APPLICANT: Soares, M. Melina

APPLICANTON: Combinations and Kits for Treating Viral Infections Using Antibod

TITLE OF INVENTION: Combinations and Kits for Treating Viral Infections Using Antibod

TITLE OF INVENTION: Aminophospholipids

FILE REFERENCE: 4001.002982

CURRENT FILING DATE: 2003-08-15

PRIOR APPLICATION NUMBER: US/10/642,060

CURRENT FILING DATE: 2003-07-15

PRIOR APPLICATION NUMBER: US 10/621,269

PRIOR APPLICATION NUMBER: 60/396,263

PRIOR FILING DATE: 2002-07-15

NUMBER OF SEQ ID NOS: 9

SOFTWARE: PatentIn version 3.1

SEQ ID NO 4

LENGTH: 144

TYPE: PRT

ORGANISM: Mus musculus

US-10-642-060-4
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US-10-642-060-4
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Search completed: April 4, 2005, 16:42:44
Job time : 16.2364 secs
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Best Local Similarity 90.0%;
Matches 9; Conservative
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46 RASQDIGSSL 55
                                                                                             1 RASQDIGNSL 10
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                                                                                                                                                                                   Score 43; DB 16; Length 144; Pred. No. 2.3; 1; Mismatches 0; Indels
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Maximum Match 100%
Listing first 45 s
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(without alignments)
266.512 Million cell update
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Copyright (c) 1993 - 2005 Compugen Ltd
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C2640551
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Description
                                                               A;Molecule type: mRNA
A;Residues: 1-106 <SHL>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;1-23/Region: framework 1
E;16-90/Domain: immunoglobulin homology <IMM>
                         F;24-34/Region: complementarity-determining F;35-49/Region: framework 2
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Best Local (
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9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                     RASQDIGSSL 15
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    complementarity-determining
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	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30
	34	<u>3</u> 5	35	ω G	35	36	36	36	36	36	36	36	36	36	36	36
	64.2	66.0	66.0	66.0	66.0	67.9	67.9	67.9	67.9	67.9	67.9	67.9	67.9	67.9	67.9	67.9
	87	422	139	130	125	730	298	128	126	122	115	115	108	108	108	108
	N	<u>, , , , , , , , , , , , , , , , , , , </u>	N	N	N	N	N	N	N	N	N	N	N	N	N	N
	PH1082	KXHUZ	S40365	PL0113	S04936	B83592	B75429	A26406	A34904	A29380	JL0080	A53276	869903	869900	S69902	S38862
1	Ig light chain V r	plasma protein Z p	Ig kappa chain - h	Ig kappa chain pre	Ig kappa chain pre	hypothetical prote	hypothetical prote	_	Ig kappa chain pre	Ig kappa chain pre	Ig kappa chain pre	Ig kappa chain V r	Ig kappa chain (cl	_	Ig kappa chain (cl	_

ALIGNMENTS

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RESULT 2 PL0260 Ig kappa chain V region (anti-DNA, DP7VK) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000 C;Accession: PL0260 C;Accession: PL0260 R;Shlomchik, M.; Mascelli, M.; Shan, H.; Raddic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A. R;Shlomchik, M.; Mascelli, M.; Shan, H.; Raddic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A. J. Exp. Med. 171, 265-297, 1990 J. Exp. Med. 171, 265-297, 1990 A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic not an autoimmune mice arise by clonal expansion and somatic not an autoimmune mice arise by clonal expansion and somatic not access to the control of th PL0261 If kappa chain V region (anti-DNA, DP15VK) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 09-May-1997 C;Accession: PL0261 C;Accession: PL0261 C;Accession: PL0261 C;Accession: PL0261 C;Accession: PL0261 A;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A J. Exp. Med. 171, 265-297, 1990 A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic A;Reference number: PL0231; MUID:90111618; PMID:2104919 A;Molecule type: mRNA A;Residues: 1-88 <SHL> C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;39-70/Region: framework 3 F;71-79/Region: complementarity-determining F;80-88/Region: framework 4 F;6-16/Region: complementarity-determining F;32-38/Region: complementarity-determining Score 43; DB Pred. No. 0.2; 1; Mismatches 2 °, Length 88 Indels 0 Gaps 0

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A;Introns: 19/1

A;Introns: 19/1

C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap C;Complex: An immunoglobulin version; such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin version; immunoglobulin homology (Skeywords: alternative initiators; heterotetramer; immunoglobulin F;1-22/Domain: signal sequence #status experimental <SIG1>
F;3-22/Domain: signal sequence #status experimental <SIG2>
F;3-22/Domain: Ig kappa chain version (MOPC 41) #status experimental <AICI)
F;3-130/Product: Ig kappa chain version (MOPC 41) #status experimental <AICI)
F;38-112/Domain: immunoglobulin homology <IMM>
F;45-110/Disulfide bonds: #status predicted
                        <sub>.</sub>Ig kappa chain
                                             RESULT 4
PL0259
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 24-Apr-1984 #sequence revision 24-Apr-1984 #text change
C;Accession: A93211; B93211; A93815; A94239; A01922; A01923
C;Accession: A93211; B93211; A93815; A94239; A01922; A01923
R;Seidman, J.G.; Max, E.E.; Leder, P.
Nature 280, 370-375, 1979
Nature 280, 370-375, 1979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     F;57-88/Region: f:
F;89-97/Region: co
F;98-106/Region:
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A;Nolecule type: protein
A;Note: Met-3 : apparently used as an alternative initiator
R;Gray, W.R.; Dreyer, W.J.; Hood, L.
Science 155, 465-467, 1967
A;Title: Mechanism of antibody synthesis: size differences be
A;Reference number: A94239; MUID:67056897; PMID:4162931
                                                                                                                                  8
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A;Residues: 23-49,'B',51-53,'LSB',57-58,'ZZ',61-62,'BZ',65-76,'B',78-108,110-130
A;Experimental source: Bence Jones protein MOPC 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Accession: B93211
A;Mccession: B93211
A;Mccession: B93211
A;Mccession: 1-117 <VK41>
A;Residues: 1-117 <VK41>
A;Residues: 1-117 <VK41>
A;Residues: 1-117 <VK41>
A;Cose-references: GB:V00804; GB:J00566; NID:g52127; PIDN:CAA24186.1; PID:g575660
A;Note: the sequences were determined from the differentiated gene MOPC 41 and the germl R;Burstein, Y; Schechter, I.
R;Burstein, Y; Schechter, I.
Proc. Natl. Acad. Sci. U.S.A. 74, 716-720, 1977
A;Title: Amino acid sequence of the NH-2-terminal extra piece segments of the precursors A;Reference number: A93815; MUID:77148916; PMID:403522
A;Accession: A93815; MUID:77148916; PMID:403522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nature 280, 370-375, 1979
A;Title: A kappa-immunoglobulin gene is formed by site-specific A;Reference number: A93211; MUID:79221900; PMID:111146
A;Accession: A93211
A;Molecule type: DNA
A;Residues: 1-130 <PC41>
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A:Accession: B93211
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                                                                                                                                                                                                                    Similarity 9; Conserv
                                                                                                                                                                      RASODIGNSL 10
                                                                                                                                RASQDIGSSL 55
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                    region
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                      (anti-DNA,
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Pred. No.
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Pred. No. 0.24;
1; Mismatches
                      DP11VK)
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0.3;
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                      mouse
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                  (fragment)
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A; Access. DNA
A; Molecule type: DNA
A; Residues: 1-95 < WAN>
C; Comment: This antibody is produced by Eps
C; Superfamily: immunoglobulin V region; imm
C; Superfamily: immunoglobulin V region; immunoglobulin
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C; Superfamil:
C; Keywords:
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PH0865
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S40330
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Best Local S
Matches 9
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Similarity 90.0%;
9; Conservation
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Similarity 90.0%;
9; Conservation
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R;Manheimer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, B.
J. Exp. Med. 174, 1639-1652, 1991
A;Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idiotype.
A;Reference number: PH0862; MUID:92078875; PMID:1660528
A;Accession: PH0865.
                                                                                                                                                                                                                                       Ig kappa chain V region (anti-DNA, I-2a) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 21-Jan-2000
C;Accession: PH0865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Residues: 1-108 <KLE>
A;Residues: 1-108 <KLE>
A;Cross-references: EMBL:X72440; NID:g441348; PIDN:CAA51108.1; PID:g441349
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eur. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi
A;Reference number: $40312; MUID:94080891; F
A;Accession: $40330
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C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
C;Accession: PL0259
R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A
J. Exp. Med. 171, 265-297, 1990
A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic n
A;Accession: PL0231; MUID: 90111618; PMID: 2104919
A;Accession: PL0259
A;Molecule type: mRNA
A;Residues: 1-106 <SHL>
A;Residues: 1-106 <SHL>
A;Residues: 1-106 <SHL>
A;Residues: 1-106 <SHL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    F;11-23/Region: framework 1
F;16-90/Domain: immunoglobulin homology <IMM>
F;24-34/Region: complementarity-determining 1
F;35-49/Region: framework 2
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Pred. No. 0.62;
0; Mismatches
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Pred. No. 0.38;
0; Mismatches
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Epstein-Barr virus-transformed immunoglobulin homology
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                            cell that bears
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RESULT 9
S19970
Ig kappa
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: $37511
R;Klein, U:; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peripheral blood A;Reference number: $37501
A;Accession: $37511
                                                                                                                                                                                                                      A;Cross-references: UNIPROT:Q65298; EMBL:X86352; NID:g807865; PIDN:CAA60124.1; PID:g8089
A;Experimental source: isolate Mexican
C;Superfamily: alfalfa mosaic virus 32.4K protein
                                                                                                                                                                                                                                                                        R;Rafael-Martin, M.; Rivera-Bustamante, R.F.
submitted to the EMBL Data Library, April 1995
A;Description: Nucleotide sequence of asparagus
A;Reference number: $54137
A;Accession: $54137
A;Molecule type: genomic RNA
A;Residues: 1-281 <RAF>
                                                                                        밁
                                                                                                                     5
                                                                                                                                                                                                                                                                                                                                                                                             probable movement protein - asparagus virus II
C;Bpecies: asparagus virus II
C;Date: 19-Mar-1997 #sequence_revision 18-Jul-1997
C;Accession: 854137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S
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A;Molecule type: mRNA
A;Residues: 1-91 <KLE>
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Best Local 9
                                                                                                                                                          Matches
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Best Local
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   kappa chain V region (M-T151) - mouse (fragment)
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                                                                                      176
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                                                                                                                      1 RASQDIGNSLT 11
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                                                                                                                                                        Similarity 7; Conservat
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                                                                                      RYSEDVGSSLT 186
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70.0%;
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81.8%;
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Pred. No. 1.4;
0; Mismatches
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Pred. No. 2.1;
                                                                                                                                                                        Score 38; DB 2; Pred. No. 6.8;
                                                                                                                                                          Mismatches
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$40353
If kappa chain V-J-C region - human
C;Species: Homo sapiens (man)
C;Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-2000
C;Accession: $40353
R;Klein, R. J Jaenichen, R.; Zachau, H.G.
Eur. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermutation.
A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Accession: $40353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;1-19/Domain: signal sequence #status predicted <SIG>F;20-114/Product: Ig kappa chain V region #status predicted <MA: F;42-107/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Straubinger, B.; Thiebe, R.; Huber, C.; Osterholzer, E.; Zachau, H.G. Biol. Chem. Hoppe-Seyler 369, 601-607, 1988
A;Title: Two unusual human immunoglobulin V-kappa genes.
A;Reference number: S00996; MUID:89114397; PMID:2852016
A;Accession: S00996
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S00996
                                                 A;Status: preliminary; translation A;Molecule type: mRNA A;Residues: 1-125 <KLE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-114 <STR>
A; Cross-references: EMB
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A;Molecule type: mRNA
A;Residuse; 1-108 <WEI>
A;Residuse; 1-108 <WEI>
A;Cross-references: EMBL:X65095; NID:g52286; PIDN:CAA46223.1; PID:g52287
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;16-90/Domain: immunoglobulin homology <IMM>
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C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change
C;Accession: S19970
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C;Date: 31-Mar-1992 #sequence_revision 31-Mar-1992 #text_change 23-Jul-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Introns: 16/1
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                           A;Cross-references: EMBL:X72463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Note: this sequence was
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Similarity 80.0%;
8; Conservation
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8; Conserv
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immunoglobulin V
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72.7%;
region; immunoglobulin homology
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Pred. No. 4.2;
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Pred. No.
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Ryfheologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; ansen, N.F.; Hughes, B.; Huizar, L.

Nature 408, 815-820, 2000

A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C. C.; Li, J.H.; Li, Y.; Liu, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali, Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.

Rizzo, M.; Rooney, T.; Kowley, D.; Sakano, H.

A;Authors: Salaberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.

A;Reference number: A86141; MUID:21016719; PMID:11130712
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A;Title: Combinatorial functions of two chimeric antibodies directed to human CD4 and on A;Reference number: PH1224; MUID:93077041; PMID:1446824
A;Accession: PH1224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein F2G19.20 [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004 C;Accession: B96512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: mRNA
A;Residues: 1-127 (WELL)
A;Residues: 1-127 (WELL)
A;Cross-references: GB:S50261; NID:g260761; PIDN:AAB24318.1; PID:g260762
A;Note: this mouse sequence was hybridized and fused with a human constant region gene
C;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig kappa chain precursor V region (M-T151) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 21-Jan-2000
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F;1-20/Domain: signal sequence #status predicted <SIG>
F;21-127/Product: Ig light chain V region #status predicted <MAT:</pre>
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A; Residues: 1-312 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   F;36-110/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                             A;Status: preliminary
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                                                                                                                                                                                                                                      A; Cross-references:
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Best Local S
Matches 8
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Best Local &
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                                                            Matches
                                                                                                                                                            position:
                                                                                  Local Similarity
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Similarity 80.0%;
8; Conservative
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       RASQDIGNSL 10
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                                                            Conservative
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                                                                                                                                                                                                                                   UNIPROT:Q9C628; GB:AE005173; NID:g11321767; PIDN:AAG34244.1; GSPDB:
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72.7%;
                                                                                    69.8%;
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Pred. No. 4.7;
1; Mismatches
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                                                                                  Score 37; DB 2;
Pred. No. 12;
                                                            Mismatches
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C;Keywords: heterotetramer; immunoglobulin v region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;1-23/Region: framework 1 F;24-34/Region: complementarity-determining 1 F;35-49/Region: framework 2 F;50-53/Region: complementarity-determining 2
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S38564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            J. Exp. Med. 174, 1639-1652, 1991
A;Title: Molecular characteristics of antibodies bearing
A;Reference number: PH0862; MUID:92078875; PMID:1660528
                                                                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-93 -KMON>
A;Cross-references: EMBL:X75105; NID:g414153; PIDN:CAA52996.1; PID:g414154
C;Superfamily: immunoglobulin
C;Keywords: immunoglobulin
C;Keywords: immunoglobulin
F;16-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig kappa chain V region (ASWU1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000
C;Accession: S38564
R;Monestier, M.; Losman, L.J.; Novick, K.E.; Aris, J.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: protein
A; Residues: 1-53 <MAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R; Manheimer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Homo sapiens (man)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 09-May-1997
C;Accession: PH0881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ig kappa chain V re N; Alternate names:
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Search completed: April Job time: 4.97125 secs
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A, Accession: S38564
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                                                                                                                                                                                                                                                                                                                                                                                                                 A; Description:
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Pred. No. !
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Pred. No. 3;
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homo sapien homo sapien homo sapien mus musculu mannheimia burkholderi

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1: uniprot_sprot:*
2: uniprot_trembl:*
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KV5K MOUSE
KV5L MOUSE
Q92539
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Q65298 mannheinia
Q65298 mannheinia
Q65298 asparagus v
Q98702 elm mottle
Q91mn5 hydrangea m
Q9c628 arabidopsis
Q6r6c8 listeria iv
Q7q448 anopheles
Q9101595 homo sapien
P01645 mus musculu
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Q6crp8 kluyveromyc
Q7z3y4 homo sapien
Q6mmj3 bdeellovibri
Q9rv58 deinococcus
Q6f613 acinetobact
Q9fv65 medicago va
Q91683 pseudomonas
Q90qv6 human immun
Q7wzp4 pseudomonas
Q87xx8 pseudomonas
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RESULT 1

KY5G_MOUSE STANDARD; PRT; 130 AA.

AC P01639; P01640;
DT 21.JUL-1986 (Rel. 01, Created)
DT 21.JUL-1986 (Rel. 01, Last sequence update)
DT 21.JUL-1986 (Rel. 04, Last annotation update)
DT 30.JUL-1986 (Rel. 04, Last annotation update)
DT 95.JUL-2004 (Rel. 44, Last annotation update)
DT 95.JUL-2004 (Rel. 40, Last annotation update)
RN SEQUENCE PROM N. A.
RX MEDLINE=79221900; PubMed=111146;
RR MEDLINE=77148916; PubMed=403522;
RN MEDLINE=77148916; PubMed=403522;
RR MEDLINE=77056897; PubMed=4162931;
RR MEDLINE=67056897; PubMed=4162931;
RR
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SMART; SM00406; IGv; 1.
SMART; SM00406; IG LIKE; 1.
PROSITE; PS50835; IG LIKE; 1.
Bence-Jones protein; Direct protein sequencing;
Immunoglobulin V region; Signal.
                                                                                                                                                                                                                                                                     -:- MISCELLANEOUS; This precursor was synthesized system directed by mRNA isolated from myeloma-i- MISCELLANEOUS; This is a Bence-Jones protein. PIR; A93211; KVMSM4.
HSSP; P01594; 1JV5.
InterPro; IPR007110; Ig-like.
InterPro; IPR0031596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=77148916; PubMed=403522;
Burstein Y., Schechter I.;
"Amino acid sequence of the NH2-terminal extra piece segments of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

MEDILINE=79221900; PubMed=111146;

MEDILINE=79221900; PubMed=111146;

Seidman J.G., Max E.E., Leder P.;

Seidman J.G., Max E.E., Leder P.;

"A kappa-immunoglobulin gene is formed by site-specific recombination without further somatic mutation.";

Nature 280:370-375(1979).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 precursors of mouse immunoglobulin lambdal-type
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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KV5I_MOUSE
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Q62HW0
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COS1O7
PRTZ HUMAN
Q8EDČ1
Q822S9
Q8XOF2
Q7YYE0
Q7YYE0
Q75JG1
Q9NDB5
KV1G_HUMAN
                         Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
      Complementarity-determining-3
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Q82289
Q7yy62
Q7yy62
Q7yy65
Q75j91
Q9npb5
P01699
P01642
Q65vu1
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kappa

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ID 29652
AC Q652
DT 01-N
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DT 01-J
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                               065298;
065298;
01-NOV-1996
01-NOV-1996
01-JUN-2003
                                                 SEQUENCE FROM N.A.

Rafael-Martin M., Rivera-Bustamante R.F.;
Rubmitted (MAY-1995) to the EMBL/GenBank/DDBJ
EMBL; X86352; CAA60124.1; -.

PIR; S54137; S54137.

InterPro; IPR002538; Bromo MP.
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Q65SW6;
25-OCT-2004
25-OCT-2004
                                                                                                                                                                                                  01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01, 01-JUN-2003 (TrEMBLrel. 24, Putative movement protein.
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SEQUENCE
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DISULFID
VARIANT
                                   SEQUENCE
                                                                                                                                                               Viruses; ssRNA positive-strand viruses, no DNA stage; Bromoviridae;
                                                                                                                                                                            Asparagus virus 2.
                                                                                                                                                                                                                                                                                                                                                                                                                            Hong S.H., Kim J.S., Lee S.Y., In Y.H., Choi S.S., Kim C.H., Jeong H., Hur C.-G., Kim J.G.; "The genome sequence of the capnophilic rumen bacta succiniciproducens."; Nat. Biotechnol. 0:0-0(2004). EMBL; AE016827; AAU37944.1; -.
                                                                                                                                         NCBI_TaxID=39681;
                                                                                                                                                        Ilarvirus
                                                                                                                                                                                         Name=ORF1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=MBEL55E;
Hong S.H., Kim J.S.,
Kim C.H., Jeong H., 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name-ompC; ORFNames=MS1337;
Mannheimia succiniciproducens MBEL55E.
Bacteria; Proteobacteria; Gammaproteob
Pasteurellaceae; Mannheimia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OmpC protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CBI_TaxID=221988;
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ses 9; Conserv
                                 PF01573; Bromo MP; 1.
NCE 281 AA; 31476 MW;
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130
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45
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90.0%;
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63.6%;
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63.6%;
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Last annotation update)
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Last annotation update)
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Pred.
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Score 38;
Pred. No.
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Pred. No. 1
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By similarity.
Missing (in 25% of the molecules).
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                                  891B345872FC2E25 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                 83DD67AF0F30BF27 CRC64;
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36;
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         Length 281;
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RESULT
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1D 098702
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OC Viru
OC Viru
OC Ilar
OC NCBI
RN [1]
RN [1]
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RN EDU
RX MEDL
RA Scott
RT rela
RL Archi
DR EMBL
DR Ffan
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Q98702;
                                                                                                                                                                               "Viruses in subgroup 2 of the genus Ilarvirus share both serological relationships and characteristics at the molecular level."; Arch. Virol. 148:2063-2075(2003).

EMBL; AF172965; AAF89807.1; -.
InterPro; IPR002538; Bromo MP.
Pfam; PF01573; Bromo MP.
Pfam; PF01573; Bromo MP.
SEQUENCE 283 AA; 31508 MW; 863CD623774291A5 CRC64;
                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
MEDLINE=22897981; PubMed=14551826;
Scott S.W., Zimmerman M.T., Ge X.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Putative movement protein 3a. Hydrangea mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UN-2001 (TrEMBLrel. 17, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9IMN5;
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EMBL; U57048; AAB97770.1; -.

InterPro; IPR002538; Bromo MP.

Pfam; PF01573; Bromo MP; 1.

SEQUENCE 283 AA; 31508 MW; 9D3B923
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Elm mottle virus.
Viruses; ssRNA positive-strand viruses, no
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01-FEB-1997 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=42812;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=97372954; PubMed=9229009; Ge X., Scott S.W., Zimmerman M.T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   larvirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         iruses; ssRNA positive-strand viruses,
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176
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                                                                                     Similarity 63. 7; Conservative
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RFSEDVGSSLT 186
                                         RASQDIGNSLT 11
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63.6%;
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Pred. No.
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Mismatches
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RESULT

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05-JUL-2004
05-JUL-2004
05-JUL-2004
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EMBL; AC083835, AAG50622.1; -.

PIR; B96512; B96512.

InterPro; IPRnoff?
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Q9C628;
01-JUN-2001
Q7QA48
Q7QA48;
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2003 (TrEMBLrel. 25, Last annotation update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protein F2G19.20.
Name=F2G19.20;
Arabidopsis thaliana (Mouse-ear cress).
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
eurosids II, Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Intraspecific Phylogeny and Lineage Group Identification Based prfA Virulence Gene Cluster of Listeria monocytogenes.";
J. Bacteriol. 186:4994-5002(2004).
EMBL; AY510073; AAR97354.1;
InterPro; IRR007752; ActA.
Pfam; PF05058; ActA; 1.
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STRAIN-NRRL 33021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Listeria ivanovii subsp. londoniensis.
Bacteria; Firmicutes; Bacillales; Listeriaceae; Listeria
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TIGRFAMs; TIGR01640; F_box_assoc_1; 1.
Hypothetical protein.
SEQUENCE 312 AA; 36256 MW; 025F1B0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=actA;
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NCE 847 AA;
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(TrEMBLrel.
                                   PRELIMINARY;
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Last sequence that the control of the cont
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Pred. No. 64;
2; Mismatches
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Pred. No. 1.9e+02;
5; Mismatches 0;
                                   PRT;
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annotation update)
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                                   4091 AA
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Best Local S
Matches 7
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InterPro; IPR001589; Actbind actnin.
InterPro; IPR001515; Calponin-like.
InterPro; IPR001449; PH.
InterPro; IPR001452; SH3.
InterPro; IPR001605; Spectrin.
InterPro; IPR001605; Spectrin.PH.
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P01595;
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SEQUENCE
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01-MAR-2004 (TrEMBLrel. 26, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation
01-MAR-2004 (TremBLrel. 26, Last annotation
AgCP14079 (Fragment).
Name=agCG48186; ORFNames=ENSANGG000000014545;
MEDLINE=73029807; PubMed=4563064;
Braun M., Leibold W., Barnikol H.U., Hilschmann N.;
"Principle of antibody structure. The primary structure of a
monoclonal kappa I-type immunoglobulin L-chain (Bence Jones parts). 3. The complete amino acid sequence and the genetic sign
                                                                                                                                                                                                                                                                                                                                                                                                                    Prodom; PD000066; SH3; 1.

PROSITE; PS00019; ACTININ_2; 1.

PROSITE; PS50021; CH; 2.

PROSITE; PS50021; CH; 2.

PROSITE; PS50003; PH_DOMAIN; 1.

PROSITE; PS50002; SH3; 1.

PROSITE; PS50002; SH3; 1.

PROSITE; PS00430; TONB_DEPENDENT_REC_1; UNKNOWN_1.
                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
MCBI_TaxID=9606;
                                                                                                                                     21-JUL-1986 (Rel. 01, Created
21-JUL-1986 (Rel. 01, Last se
05-JUL-2004 (Rel. 44, Last an
05-JUL-2007 (Rel. 47) (Rejon Bi.
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Pfam; PF00159; PH; 1.
Pfam; PF00018; SH3 1; 1.
Pfam; PF00413; Spectrin; 31.
PRINTS; PR00683; SPECTRINPH.
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GO; GO:0003779; F:a
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Anopheles.
                                                                                                                             Homo sapiens (Human)
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4091 AA;
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A; 469663
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; Spectrin_PH.
; TONB_Box_N.
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Pred. No. 9.9e+02;
3; Mismatches 0
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|on update
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variability principles for the mechanism of antibody

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RESULT

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Matches 8
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Bence-Jones protein; Direct pro
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMARP; SM004406; IGy; 1.
PROSITE; PS50835; IG_LIKE; 1.
Direct protein sequencing; Immunoglobulin V
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NON TER
SEQUENCE
                                                                                                                                                                                                                                                 PIR; A01927; KVMSAR.
HSSP; P01594; 1JV5.
                                                                                                                                                                                                                                                                                                          "Complete amino acid sequence of light chain variable regions derived from five monoclonal anti-p-azophenylarsonate antibodies differing with respect to a crossreactive didotype.";

Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
                                                                                                                                                                                                                                                                                                                                                                                              Siegelman M.,
                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=A/J;
MEDLINE=82150934; PubMed=6801658;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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21-JUL-1986 (Rel. 01,
05-JUL-2004 (Rel. 44,
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-!- MISCELLANEOUS: The C re-
-!- MISCELLANEOUS: This is
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HSSP; P01607; 1BWW.
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    Physiol. Chem. 353:1284-1306(1972).
    The C region of this chain has the INV (3) marker.
    This is a Bence-Jones protein.

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Pred. No.
1; Mismatc
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Complementarity-determining-3
                                   Framework-2.
Complementarity-determining-2
                                                                              Complementarity-determining-1.
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RESULT 12

KV5M MOUSE

ID KV5M MOUSE

AC P01646;

DT 21-UUL-1986

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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
05-JUL-2004 (Rel. 44, Last annotation
                  Ig kappa chain V-V region HP 123E6.
Mus musculus (Mouse).
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SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LI
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"Complete amino acid sequence of light chain variable reg
from five monoclonal anti-p-azophenylarsonate antibodies
with respect to a crossreactive idiotype.";
Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683 (1981)
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21-JUL-1986 (Rel. 01,
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HSSP; P01607; 1REI.
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InterPro; IPR003596; Ig_v.
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05-JUL-2004 (Rel. 4
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981)
                                                                                                                                                                                                                                                                                                                                                               from five monoclonal anti-p-azophenylarsonate ant with respect to a crossreactive idiotype.", Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
-I- MISCELLANEOUS: Anti-arsonate hybridoma protei
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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InterPro; IPR003596; Ig_v.
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RESULT 14
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 Q925S9;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Immunog.obulin light chain (Fragment).
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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation updat
                                                                                                                                                                                                                                                                             Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LI
                                                                                                                                                                                                                                                                                                                                            Siegelman M., Capra J.D.;

"Complete amino acid sequence of light chain variable regions derived from five monoclonal anti-p-azophenylarsonate antibodies differing with respect to a crossreactive idiotype.";

Proc. Natl. Acad. Sci. U.S.A. 78:7679-7683(1981).
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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InterPro; IPR003596; Ig_v.
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Search completed: April 4, 2005, 15:56:08 Job time : 20.5911 secs
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RC STRAIN=B9J06687; PubMed=10380019;

REDLINE=99306687; PubMed=10380019;

RX MEDLINE=99306687; PubMed=10380019;

RX Tilpathi P.K., Qin H., Bhattacharya-Chatterjee M., Ceriani R.L.,

RA Tilpathi P.K., Qin H., Bhattacharya-Chatterjee M., Ceriani R.L.,

RA Tilpathi P.K., Qin H., Bhattacharya-Chatterjee M., Ceriani R.L.,

RAD TILPATHIC P.K.,

ROON TER 127 127

SEQUENCE 127 AA; 13794 MW; 13F61BEBBB981FA5 CRC64;
                                                                                                                                                                                                              Query Match 67.9%; Score 36; DB 2; Length 127; Best Local Similarity 80.0%; Pred. No. 40; Matches 8; Conservative 1; Mismatches 1; Indels
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
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Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match I	Length	B	Ħ	Description
1	31	100.0	7	2	AAW70945	Aaw70945 CDR2 of t
N	31	•	7	w	AAY44590	Mouse
w	31	•	7	ð	ABR55316	Comple
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6	31	•	30	σ	ABR55336	Amino
7	31	100.0	83	8	ADM91687	Adm91687 Lawsonia
8	31	100.0	90	ผ	AAR80078	Aar80078 Mouse der
9	31	100.0	92	N	AAW95480	Aaw:5480 Mouse der
10	31	100.0	92	N	AAW95476	Aaw95476 Mouse der
11	31	100.0	95	N	AAW70954	Aaw70954 Light cha
12	31	100.0	97	v	ABP33764	
13	31	100.0	99	8	ADM78077	Adm78077 Human SJA
14	31	100.0	107	N	AAW27121	Aaw27121 Murine an
15	31	100.0	107	w	AAY90824	52009
16	31	100.0	108	ผ	AAR21310	Aar21310 Light cha
17	31	100.0	108	w	AAY44587	Aay44587 Mouse ant
18	31	100.0	108	σ	ABR55325	
19	31	100.0	108	δ	ABR55319	Abr55319 Variable
20	31	100.0	108	0	ABR55331	
21	31	100.0	108	o	ABR55329	Abr55329 Amino aci
22	31	٠	108	0	ABR55330	0 Amino
23	31	100.0	108	œ	ADI26702	Mouse
24	31	100.0	108	ω	AD060445	Ado60445 Humanised
25	31	100.0	108	ω	ADO60443	Ado60443 Mouse ant

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AAW70945 standard; peptide; 7 ₹

ALIGNMENTS

AAW70945;

14-OCT-1998 (first entry)

CDR2 of the light chain of monoclonal antibody RS-255

Pathogenic virus; tropism; mucosa; CDR region; monoclonal antibody; respiratory syncytial virus; RSV; VP6 protein; rota virus; RV; viral infection; inhibit, fusion; protection; transcription; antiviral agent; prophylaxis; diagnosis; infection; contamination.

Synthetic

17-JUL-1998 FR2758331-A1

14-JAN-1997; 97FR-00000300.

14-JAN-1997; 97FR-00000300

(UYBO-) UNIV BOURGOGNE.

WPI; 1998-390320/34.

peptide(s) recognising viral epitope with tropism to mucosa - useful, e.g. diagnosing, preventing and treating viral infection(s).

Claim 12; Page 39; 51pp; French.

RRESULT 1
AAW7094
ID AAW77
XX AAW77
XX AAW77
XX AAW77
XX CDR2
XX CDR2
XX Path
XX W resp
XX VITA
OS Mus
XX Synt
OS Mus
XX I1-J
XX I1-J
XX I1-J
XX IVYB
XX IVYB AAW70905-46 represent peptide sequences that can recognise, by antigenantibody type reactions, at least I epitope of a pathogenic virus having tropism for the mucosa. AAW70905-16 and AAW70929-46 are analogous to CDR regions of monoclonal antibodies specific for respiratory syncytial virus (RSV). AAW70917-28 are analogous to CDR regions of monoclonal antibodies specific for site III or IV of the VP6 protein of rota virus (RV). The peptides can neutralise viral infections and may also inhibit fusion between infected and uninfected cells or cells and viruses. They provide passive or active protection and/or inhibit transcription of the virus, so are useful as antiviral agents or for prophylaxis, in human or veterinary medicine. The peptides can be labelled and used to diagnose infection or contamination by the virus. The peptides are particularly directed against RSV or RS but may also be used against papilloma, adeno,

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RESULT 2
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                                                                                                                                                                                                                                                                The present protein sequence is a complementarity-determining region (CDR) of mouse anti-Interleukin-18 antibody light chain variable region (VL) encoded by cDNA derived from hybridoma #125-2H. The nucleotide sequence can be used in the production of recombinant monoclonal antibody #125-2HmAb, which is capable of neutralising biological activities of interleukin-18. The antibody has antiinflammatory, immunosuppressive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New artificially produced peptide for neutralizing biological activity of interleukin-18, useful for treating and preventing immunopathies, inflammatory disorders and autoimmune diseases.
                                                                                                                  Sequence 7 AA;
                                                                                                                                                                  leucocytopoietic, antialgic, antipyretic, antiallergic and hepatotropic activity and can be used for prevention and treatment of autoimmune diseases, immunopathies and inflammatory disorders caused by excessive immunoreaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 7; Page 18; 32pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-118341/11.
N-PSDB; AAZ49537.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nishida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JUN-1998;
12-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complementarity-determining region; CDR; hybridoma #125-2H; mouse monoclonal antibody #125-2HmAb; interleukin-18; antiinflammatory; immunosuppressive; leucocytopoietic; antialgic; antipyretic; antiallergic; hepatotropic; treatment; autoimmune disease; immuno; inflammatory; disease; immuno;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti-Interleukin-18 antibody; IL-18; light chain variable region; VL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY44590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY44590 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HAYB ) HAYASHIBARA SEIBUTSU KAGAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse anti-IL-18 antibody VL complementarity-determining region-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entero, polio, influenza or immune deficiency viruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammatory
     Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATSSLDS
        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Okura T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; ilarity 100.0%; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder; immunoreaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98JP-00177580.
98JP-00289044.
98JP-00365023.
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                            100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tanimoto T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
     0,
Score 31; DB 3; Pred. No. 1.8e+06; Nismatches 0;
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Pred. No. 1.8
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kurimoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2; ]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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  Indels
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RESULT 3
ABR55316
AD ABR55316
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AX ADR1
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RESULT 4
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes an antibody molecule which has specificity for human kinase insert domain-containing receptor (KDR). The antibody comprises complementarity determining regions (CDRs) from the heavy and light chain variable domains of the murine monoclonal antibody VR165. This antibody is specific for human KDR. The antibody of the invention blocks vascular endothelial growth factor (VEGF) binding to KDR. Antibodies of the invention are useful for treating a pathology in which VEGF and/or KDR are implicated, and in the manufacture of a medicament for the treatment of a pathology in which VEGF and/or KDR are implicated. This includes inflammation, psoriasis, rheumatoid arthritis, and tumour growth or metastasis. ABR55315-17 represent the CDRs of the light chain variable region of VR165 antibody. They are used to produce antibodies of
                                                                                             ADM78136;
                                                                                                                                                       ADM78136 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antibody molecules having specificity for human kinase insert domain-containing receptors, useful for treating inflammation, psor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 2; Fig 1; 57pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-2001; 2001GB-00024317.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-OCT-2002; 2002WO-GB004619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Complementarity determining region of light chain of antibody VR165.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CELL-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 100 7; Conservative
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                                                                                                                                                                                                                                                                                                               ATSSLDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tickle
                                                                                                                                                 peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 31; DE 100.0%; Pred. No. 1.8 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumor growth and metastasis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                1.8e+06;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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01-JUL-2004

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel monoclonal antibody specific to adenylate kinase isozyme 3, comprising 4 or more of 6 complementarity determining regions (CDRs) of a sequence having 6-18 amino acids. The invention further relates to: a composition comprising the monoclonal antibody for detecting a heart disease marker AK3 a kit comprising the monoclonal antibody for the diagnosis of heart disease; and a method of detecting a heart disease marker AK3. The monoclonal antibody is useful for preparing a composition for detecting a heart disease marker AK3. The monoclonal antibody reduces false positive results of the conventional biochemical markers. This sequence represents a human antibody variable light chain CDR peptide region of the invention.
                                                                                                     neuroprotective; targeted therapy; humanised antibody; human insulin receptor; HIR; heavy chain; constant region; human tregion; complementarity determining region; CDR; framework; neuropharmaceutical agent; blood brain barrier; avidin-bictin linkage; mouse; antibody 83-14; Mab 83-14; light chain variable region; VL; CDR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New monoclonal antibody specific to human mitochondrial adenylate kinase isozyme 3, useful for preparing a composition for detecting a heart disease marker AK3.
                                               US2004101904-A1
                                                                                                                                                                                                    Mouse antibody 83-14 light chain CDR2.
                                                                                                                                                                                                                                    12-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                  ADO60479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                monoclonal antibody; adenylate kinase isozyme 3; complementarity determining region; CDR; heart disease; maxker AK3; biochemical; human; variable; light chain.
                27-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-316088/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-SEP-2002; 2002KR-00059211.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human SJB3-86 antibody variable light chain CDR2 peptide region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KIMH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                              1 ATSSLDS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 100 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 99; 126pp; English.
                                                                                                                                                                                                                                                                                                  standard; peptide; 7 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2003WO-KR001979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 31; DB 8;
Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 7;
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RESULT 6
ABR55336
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC variable region and a second CDR located between the first and third CC CDRs. The framework regions comprise a first frame work region located adjacent to the amino end of the first CDR, a third framework region CC located between the first and second CDR, a third frame work region CC located between the second and third CDR and a fourth frame work region CC located adjacent to the carboxyl end of the third CDRs, where the CC framework regions have been humanized, and a light chain of aminoacids CC compatible with the heavy chain. (I) is useful for increasing the ability CC involves linking the neuropharmaceutical agent to (I) by an avidin-biotin CC linkage or by genetic fusion. (I) is useful for targeting a CC compartical agent to (I). A preparation (II) comprising (I) is useful for intravenously administering a pharmaceutical agent to a human patient CC is useful in combination with drug and diagnostic agent to treat human CC is useful in combination with drug and diagnostic agent to treat human CC carrians in vivo. (I) is useful as Trojan horse to transport to the amino CC determining region 2 (CDRR). Showing the preferred sequence of the CDR constraints across blood brain barrier.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                Synthetic
                                                      complementarity determining region; CDR; VEGF; inflammation; psoriasis;
                                                        Antibody; kinase insert domain-containing receptor; KDR; antibody VR165; complementarity determining region; vascular endothelial growth factor; CDR; VEGF; inflammation; psoriasis; rheumatoid arthritis; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chain of amino acide comprising a constant region and a variable region where the variable region has an amino end and a carboxyl end and comprises complementarity determining regions (CDR) interspersed between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transport neuropharmaceutical agents, has heavy and light amino acid chain having complementarity determining regions interspersed between humanized framework regions.
                                                                                                                                      Amino acid sequence of cassette used to construct a light chain graft.
                                                                                                                                                                              29-JUL-2003
                                                                                                                                                                                                                      ABR55336;
                                                                                                                                                                                                                                                          ABR55336 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  within the humanised antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   determining region 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 framework regions, the CDR comprises a first CDR located at the amino of the variable region, a third CDR located at the carboxyl end of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a humanised antibody (I) that binds with human insulin receptor (HIR. The humanised antibody comprises a h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; SEQ ID NO 40; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Humanized murine antibody that bind to human insulin receptor useful transport neuropharmaceutical agents, has heavy and light amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                    ATSSLDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CDR2), showing the preferred sequence
                                                                                                                                                                                                                                                              30
                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 31; DB 8; Le
Pred. No. 1.8e+06;
Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of the CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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WPI; 2004-340902/31.

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Best Local S
Matches 7
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                                                                                                                                                                                                                       Lawsonia intracellularis; infection; antibacterial; immunotherapy,
                    Kapur V,
                                                                      04-OCT-2002; 2002US-0416395P.
                                                                                                   01-OCT-2003;
                                                                                                                                                           WO2004033631-A2
                                                                                                                                                                                   Lawsonia intracellularis.
                                                                                                                                                                                                              proliferative
                                                                                                                                                                                                                                                                                15-JUL-2004
                                                                                                                                                                                                                                                                                                             ADM91687;
                                               (MINU ) UNIV MINNESOTA.
                                                                                                                                                                                                                                                  Lawsonia intracellularis specific amino acid sequence SEQ ID NO:88
                                                                                                                                                                                                                                                                                                                                     ADM91687 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel antibody molecules having specificity for human kinase insert domain-containing receptors, useful for treating inflammation, psoriasis, rheumatoid arthritis, tumor growth and metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Fig 13; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-441133/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-OCT-2002; 2002WO-GB004619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-APR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACC70039.
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                    Gebhart CJ
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                                                                                                   2003WO-US031318
                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                              enteropathy.
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                                                                                                                                                                                                                                                                                                                                     protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 31; DB Pred. No. 7; O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
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RESULT 8
AAR80078
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Best Local S
Matches 7
Region
                                Region
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                                                                                                                                                                                            Key
                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                              Light chain; RT3; murine; catalytic antibody; bacteriophage; pattern
                                                                                                                                                                                                                                                                                             Mouse derived light chain RT3 phage antibody pattern A.
                                                                                                                                                                                                                                                                                                                                                                                                    AAR80078 standard; protein; 90
                                                                                                    Region
                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                     22-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid that generates an amplification product from L
intracellularis nucleic acid using an appropriate second nuclei
molecule, useful for treating and preventing L. intracellularis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 88; 87pp; English.
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7; Conserv
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                                                                                    /note=
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                                                                                                                                                                                        Location/Qualifiers
                note=
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 . 88
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100.0%; E
tive 0;
                "framework region
                                                 "complementarity determining region
                                                                                  "framework region
                                                                                                                     "complementarity determining region
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Pred. No. 21;
D; Mismatches
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21;
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WO9527045-A1

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Best Local
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                                                           (MCCA/) MCCAFFERTY J.
(CHIS/) CHISWELL D.
(DARS/) DARSLEY M J.
(TITM/) TITMAS R C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT04625 encodes AAR80078 mouse derived light chain RT3 phage antibody. The DNA was used in the prepn. of catalytic antibody (CA) groducing bacteriophage. The CAs can be used to activate/deactivate a biological function in an animal by enhancing the rate of cleavage, or formation of a specific bond within a mol. in vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                prodrug; scFV.
                                                                                                                                                                                                                                                                                                                                                                                                   Catalytic; antibody; phage display; immunising; phage expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW95480;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW95480 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Production of catalytic antibodies displayed on phage - gene library of antibody-derived domains and expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-OCT-1995
                                                                                                                                                                      22-JAN-1993;
                                                                                                                                                                                                              14-JUL-1994;
                                                                                                                                                                                                                                                        05-JAN-1999
                                                                                                                                                                                                                                                                                                US5855885-A
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse derived RT3 phage antibody light chain pattern C genetic sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 90 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 9; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; AAT04625
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAR-1994;
                   (MART/)
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               CHISWELL D.
DARSLEY M J.
TITMAS R C.
MARTIN M T.
KENTEN J H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 7, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATSSLDS 47
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, Martin MT, Titmas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
llarity 100.0%;
Conservative (
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                                                                                                                                                                      93US-00007684.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chiswell D, Darsley MJ,
itmas RC, Williams RO;
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Pred. No. 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          y generating of in phage
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note= "complementarity determining region 3"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to methods for producing catalytic antibodies CC displayed on a phage. The method comprises: (a) generating a gene library CC of antibody-derived domains; (b) inserting coding for the domains into a CC phage expression vector; and (c) isolating the catalytic antibodies. The phage expression vector incorporates a histidine peptide in tandem with a CC myc peptide. The catalytic antibodies can be isolated by preparing an CC expression vector incorporates a histidine peptide in tandem with a CC library of VH and VL domains from the immunised animal; cloning the VH CC and VL domains into a phage expression vector to generate phage display CC antibodies; selecting phage display antibodies which bind specifically to the antigen; gereaning the selected phage display antibodies for CC catalytic activity to substrate; and isolating the catalytic antibodies, which can be used for in corporates a histidine peptide in CC cantibodies, which can be used for in vivo activation of a produce catalytic contibodies, which can be used for in vivo activation of a produce The pattern CC catalytic can bequence represents a genetic sequence of light chain PCR pattern CC crom mouse derived RT3 phage antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                            (MCCA/)
(CHIS/)
(DARS/)
(TITM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 92
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                                                                                                                                                                    22-JAN-1993;
                                                                                                                                                                                                                                                                               05-JAN-1999
                                                                                                                                                                                                                                                                                                                                      US5855885-A
                                                                                                                                                                                                                                                                                                                                                                                           Mus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                               prodrug;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Catalytic; antibody; phage display; immunising; phage expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse derived RT3 phage antibody light chain pattern A genetic sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAR-1999
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N-PSDB; AAX00879.
                                                                                                                                                                                                                         14-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (FITZ/) FITZGERALD K. (WILL/) WILLIAMS R O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2
) MCCAFFERTY J.
) CHISWELL D.
) DARSLEY M J.
) TITMAS R C.
) MARTIN M T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATSSLDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            BCFV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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niswell D,
                                                                                                                                                                    93US-00007684
                                                                                                                                                                                                                           94US-00273146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MJ, William, Mccafferty J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Williams RO, Smr...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 31; DB
Pred. No. 23;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ۰,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 domains inserting
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RESULT 11
AAW70954
ID AAW70
XX AW70
AC AAW70
XX Light
XX Pathc
KW Pathc
KW respi
KW viral
KW viral
KW viral
KW antiv
XX Mus s
PN FR275
PN 17-J1
XX 14-J1
XX 14-J1
PR 14-J1
PR 14-J1
PR 14-J1
PR 14-J1
PR 14-J1
PR 14-J1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC displayed on a phage. The method comprises: (a) generating a gene library CC of antibody-derived domains; (b) inserting coding for the domains into a CC phage expression vector; and (c) isolating the catalytic antibodies. The CC myc peptide. The catalytic antibodies can be isolated by preparing an CC antigen; optionally immunising an animal with the antigen; generating a CC library of VH and VL domains from the immunised animal; cloning the VH CC and VL domains into a phage expression vector to generate phage display antibodies; selecting phage display antibodies which bind specifically to CC antibodies; selecting the selected phage display antibodies for CC catalytic activity to substrate; and isolating the catalytic antibodies, which can be used for in vivo activities peptide in CC candem with a myc peptide. The processes are used to produce catalytic antibodies, which can be used for in vivo activation of a prodrug. The present sequence represents a genetic sequence of light chain pattern A CC from mouse derived RT3 phage antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                             Pathogenic virus; tropism; mucosa; CDR region; monoclonal antibody; respiratory syncytial virus; RSV; VP6 protein; rota virus; RV; viral infection; inhibit; fusion; protection; transcription; antiviral agent; prophylaxis; diagnosis; infection; contamination.
                                          14-JAN-1997;
              (UYBO-) UNIV
                                                                                                                17-JUL-1998.
                                                                                                                                              FR2758331-A1.
                                                                                                                                                                                                                                                                                               Light chain of monoclonal antibody RS-255.
                                                                                                                                                                                                                                                                                                                                      14-OCT-1998
                                                                            14-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                      AAW70954 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 92 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Production of catalytic antibodies displayed on bacteriophages - comprises generating a gene library of antibody-derived domains inserting coding into a phage expression vector and isolating the catalytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fitzgerald K, Darrer TH, Chiswell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KENT/)
(SMIT/)
(FITZ/)
(WILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to methods for producing catalytic antibodies displayed on a phage. The method comprises: (a) generating a gene l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Fig 9A-F; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibodies.
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DB; AAX00875.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42
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) SMITH R.
) FITZGERALD K.
) WILLIAMS R O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATSSLDS 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
              BOURGOGNE
                                                                                                                                                                                                                                                                                                                                  (first entry)
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hiswell D,
                                            97FR-00000300.
                                                                            97FR-00000300.
                                                                                                                                                                                                                                                                                                                                                                                                    protein; 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MJ, Williams RO, Smith R, Mccafferty J, Titmas RC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 31;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Martin MT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Leach MD,

Shimkets RA;

24-MAY-2000; 2000US-0206690P

24-MAY-2001; 2001WO-US017076

29-NOV-2001. WO200190366-A2

(CURA-) CURAGEN CORP

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RESULT 12
ABP33764
ID ABP33764
ID ABP33764
ABP33764
ABP33764
AC ABP33
XX ABP33764
AC ABP33764
AC ABP33764
AC ABP33764
AC ABP3764
AC ABP376
AC ABP3
                                                                                                                                                                                                                                                                                                                                                                                                                                                       KW Human; ORF; open reading frame; ORFX; drug screening; diagnosis;
KW disease monitoring; cytokine; cell proliferation; cell differentiation;
KW immune modulation; haematopoiesis regulation; tissue growth;
angiogenesis; activin; inhibin; chemotactic; chemokinetic; haemostatic;
KW thrombolytic; tumour inhibition; bodily characteristic; fertility;
KW behaviour; cancer; proliferative disorder; neurological disorder;
KW cardiovascular disease; immune system disorder; organ transplantation;
KW tissue growth disorder; tissue regeneration disorder; diabetes mellitus;
KW typothyroidism; cholesterol ester storage disease; infection; vulnerary;
KW vasotropic; antipsoriatic; antidiabetic; cytostatic; nootropic;
KW neuroprotective; antiatherosclerotic; anticoagulant; thrombolytic;
KW cardiant; hypotensive; antithyroid; antipacterial functions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human transport protein-like ORF2737 protein, SEQ ID NO:5474.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP33764;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP33764 standard; protein; 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 95 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   deficiency viruses
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptide(s) recognising viral for, e.g. diagnosing, preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-390320/34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATSSLDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     epitope with tropism to mucosa - useful and treating viral infection(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2;
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RESULT 13
ADM78077
ID ADM78
XX ADM78
AC ADM78
XX 01-JU
DT 01-JU
XX MONOG
KW MONOG
KW COMp]
KW bioch
XX Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC range of biological activities, such as cytokine, cell proliferation, cc call differentiation, immune modulation, haematopoiesis regulation, cc tissue growth, angiogenesis activity or inhibin activity, chemokactic/ cc chemokinetic activity, haemastatic activity, thrombolytic activity, chemokactic/ cc receptor/ligand, antiinflammatory activity, thrombolytic activity, companies, fertility and behaviour of the determination of oblody characteristics, fertility and behaviour of cancers, cc on bodily characteristics, fertility and behaviour of cancers, companies, companies, and antibodies may be used in the treatment of cancers, companies, companies, immune system disorders and benign tumours, cc ordarovascular diseases, immune system disorders, disorders related to crystal transplantation, disorders of tissue growth and regeneration, cc diseases such as disbetes mellitus, hypothyroidism, and cholesterol ester storage disease, and infectious diseases caused by viral, bacterial, cc fungal and other pathogens. ORFX nucleic acids may also be used as a cource of primers and probes, in the detection of ORFX genomic sequences or transcripts, in the identification and cloning of homologous cancers, in the detection and cloning of homologous or transcripts, in the disances, and in forensic biology. The ORFX protein, and in drug screening. The ORFX proteins may also be used as a companies, and in drug screening. The ORFX proteins may also be used as companies, treatment and monitoring of ORFX associated disances.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     designated ORF (open reading frame) 1-4534, and sequences ABN75054-BBN7587 represent cDNAs encoding them. The invention also encompasses polypeptides at least 80% identical to the ORF1-ORF4534 (collectively referred to as ORFX) proteins, polynucleotides at least 85% identical to the ORFX nucleic acid sequences, vectors and host cells comprising ORFX polynucleotides, the recombinant production of ORFX proteins, antibodies specific for ORFX proteins, methods of detecting ORFX polynucleotides and polypeptides, methods of screening for modulators of ORFX expression or activity, and methods of screening individuals for a predisposition to an ORFX-sssociated disorder. The ORFX proteins of the invention have a wide
                                                               complementarity biochemical; hun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human polypeptides and polynucleotides useful for diagnosing, preventing and treating cardiovascular disease, neurodegenerative, hyperproliferative disorders and disorders related to organ
                       Homo sapiens
                                                                                                              monoclonal antibody; adenylate kinase isozyme 3;
                                                                                                                                                         Human SJA3-86 antibody variable light chain region,
                                                                                                                                                                                                     01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                             ADM78077 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 97 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences ABP31028-ABP35561 represent 4534 novel human proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 10; Page 1619-1620; 2508pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-106200/14.
N-PSDB; ABN77790.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                           52
                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ATSSLDS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                               lty determining region; CDR; heart disease; marker AK3; human; variable; light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and monitoring of ORFX-associated diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                               99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31; DB
Pred. No. 25;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5; Length 97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               adenylate kinase isozyme 3, comprising 4 or more of 6 complementarity determining regions (CDRs) of a sequence having 6-18 amino acids. The invention further relates to: a composition comprising the monoclonal antibody for detecting a heart disease marker AK3; a kit comprising the monoclonal antibody for the diagnosis of heart disease; and a method of detecting a heart disease marker AK3. The monoclonal antibody is useful for preparing a composition for detecting a heart disease marker AK3. The monoclonal antibody reduces false positive results of the conventional biochemical markers. This sequence represents a human variable light chain region protein of the invention.
                                                                                                                                                Region
                                                                                                                                                                 Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New monoclonal antibody specific to human mitochondrial adenylate kinase isozyme 3, useful for preparing a composition for detecting a heart disease marker AK3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Xi.m
              WO9722699-A2.
                                                                  Region
                                                                                                         Region
                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                   Monoclonal antibody 11D10; anti-idiotype antibody; mucin; human milk fat globule; HMFG; tumour; breast cancer; vacc
                                                                                                                                                                                                                                                             Murine antibody light chain variable region consensus
                                                                                                                                                                                                                                                                                         25-MAR-2003
04-JAN-1998
                                                                                                                                                                                                                                                                                                                                   AAW27121;
                                                                                                                                                                                                                                                                                                                                                           AAW27121 standard; protein; 107 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 99 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel monoclonal antibody specific to adenylate kinase isozyme 3, comprising 4 or more of 6 complements
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-SEP-2002; 2002KR-00059211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-SEP-2003; 2003WO-KR001979.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2004029094-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; SEQ ID NO 40; 126pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KIMH/) KIM H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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DB; ADM78076.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ATSSLDS 47
                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                Location/Qualifiers
                                                                                               /label= CDR2
                                                                                                                       /note= "complementarity determining region 1"
                                                      'label= CDR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                      "complementarity determining region 3"
                                                                               "complementarity determining
                                                                                                                                       CDR1
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Pred. No. 25;
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RESULT 15
AAY90824
ID AAY90820
XX AAY90
AC AAY90
AC AAY90
XX 29-AU
XX Antig
KW Antig
KW antib
KW toxin
XX Homo
XX Homo
XX 155-AF
PR 08-FE
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Best Local Similarity
Matches 7; Conserv
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26-JAN-1996;
13-DEC-1996;
                                          08-FEB-1984;
11-JAN-1985;
21-MAR-1986;
08-MAY-1988;
11-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This polypeptide sequence comprises a consensus sequence of murine light chain variable regions (VL) selected on the basis of identity to the VL region (see W27119) of monoclonal anti-iduotype antibody 11D10. The sequences were obtained from a GenBank database search A VH consensus (W27122) was also produced. 11D10 has at least 18 departures from the consensus sequences (7 in the light chain and 11 in the heavy chain). 8 occur within CDRs and 10 outside CDRs. 11D10 polypeptides and polymucleotides can be used in vaccines and pharmaceutical compositions for the treatment of human milk fat globule-associated diseases such as breast cancer. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monoclonal anti-idiotype antibody 11D10 - human milk fat globule disease associated
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                                                                                                                                                                                                                                      Antigen binding site; immunoglobulin; cancer antigen; immunological; antibody; tumour; human; mucin; cancer; cytostatic; hybridoma; specific binding assay; affinity purification; drug targeting; toxin targeting; imaging; genetic; therapeutic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Fig 26C; 130pp; English.
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                                                                                                                                                                                   US6054561-A.
                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                            520C9 hybridoma VL domain SEQ ID NO:26
                                                                                                                                                                                                                                                                                                                                      29-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                          AAY90824 standard; protein; 107 AA.
                                                                                                                              07-JUN-1995;
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96US-00591965.
96US-00766350.
                                          84US-00577976.
85US-00690750.
86US-00842476.
88US-00190778.
94US-00288981.
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                                                                                                                              95US-00483749
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Pred. No. 27;
0; Mismatches
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27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 elicits immune response against tumours, especially breast
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Search completed: April Job time: 15.9776 secs
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Matches 7
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                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 13; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                       treating cancer.
                                                                                                                                                                                                                                                                                                                                                                    Monoclonal antibody capable of binding useful for affinity purification, drug
                                                                                                                                                                      Sequence 107
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7; Conserv
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Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/IS06_NEW_PUB.pep:*
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11 US-09-864-408A-5474
12 US-09-924-099-1
14 US-10-010-729-45
16 US-10-307-276B-4
16 US-10-307-276B-6
16 US-10-30-62-267
16 US-10-803-653-267
17 US-10-803-653-267
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Sequence 4, Appli
Sequence 40, Appl
Sequence 5474, Ap
Sequence 5474, Ap
Sequence 45, Appli
Sequence 4, Appli
Sequence 6, Appli
Sequence 67, App
Sequence 267, App
Sequence 74, Appl
Sequence 74, Appl
Sequence 74, Appl
Sequence 1, Appli
Sequence 1, Appli
Sequence 35, Appli
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31 100.0 130 14 US-10-010-729-71 Sequence 71, Appl 31 100.0 144 16 US-10-642-120-4 Sequence 4, Appli 31 100.0 144 16 US-10-642-122-4 Sequence 4, Appli 31 100.0 144 16 US-10-621-269-4 Sequence 4, Appli 31 100.0 144 16 US-10-621-269-4 Sequence 4, Appli 31 100.0 144 16 US-10-620-850-4 Sequence 4, Appli 31 100.0 144 17 US-10-620-850-4 Sequence 4, Appli 31 100.0 144 17 US-10-642-100-4 Sequence 4, Appli 31 100.0 144 17 US-10-642-100-4 Sequence 4, Appli 31 100.0 144 17 US-10-642-100-4 Sequence 4, Appli 31 100.0 234 15 US-10-3698-6340 Sequence 6, Appli 31 100.0 237 9 US-09-924-099-10 Sequence 9, Appli 31 100.0 237 9 US-09-924-099-10 Sequence 6, Appli 31 100.0 243 9 US-09-924-099-10 Sequence 6, Appli 31 100.0 243 17 US-10-868-55 Sequence 6, Appli 31 100.0 252 15 US-10-239-656-57 Sequence 6, Appli 31 100.0 257 15 US-10-239-656-57 Sequence 6, Appli 31 100.0 276 9 US-09-766-543-12 Sequence 10, Appli 31 100.0 276 9 US-09-766-543-12 Sequence 10, Appli 31 100.0 276 9 US-09-766-543-12 Sequence 10, Appli 31 100.0 276 9 US-09-766-543-12 Sequence 12, Appli 31 100.0 276 9 US-09-766-543-12 Sequence 12, Appli 31 100.0 276 9 US-09-766-543-12 Sequence 12, Appli 31 100.0 567 15 US-10-369-493-1974 Sequence 10, Appli 31 100.0 567 15 US-10-369-493-1961 Sequence 10, Appli 51 US-10-369-493-1961 Sequence 10, Appli 51 US-10-369-493-1961 Sequence 10, Appl
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ALIGNMENTS

RESULT 1 US-09-924-099-4

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                                                                                                                                                                                                          SEQ ID NO 4
LENGTH: 7
TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                Query Match
Best Local :
                                                                                            Matches
                                                                                                                                                                                                                                                                                                                     APPLICANT: TANIMOTO, Taddao
APPLICANT: KURIMOTO, Masashi
TITLE OF INVENTION: PEPTIDE
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/09/924,099
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/338,511
PRIOR FILING DATE: EARLIER FILING DATE: 1999-06-23
PRIOR PILING DATE: EARLIER FILING DATE: 1999-06-23
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: JP 289,044/98
PRIOR FILING DATE: EARLIER FILING DATE: 1998-10-12
PRIOR PILING DATE: EARLIER FILING DATE: 1998-10-12
PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: NISHIDA, Yoshihiro APPLICANT: OKURA, Takanori
                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 33
                                           |-4
                                                                                       Similarity 7; Conserv
                                           ATSSLDS
                                                                                            Conservative
                                                                                       100.0%; Score 31; DB 9; 100.0%; Pred. No. 1.3e+06; vative 0; Mismatches 0;
                                                                                                                                   Length 7;
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RESULT

US-10-307-276B-40

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Query Match
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                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Leach, Martin D.
APPLICANT: Shimkets, Richard A.
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: No. US20040009474A1el Human Polynucleotides and Polypeptides EncoFILE REFERENCE: 21402-012
                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5474
LENGTH: 97
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5474, Application US/09864408A Publication No. US20040009474A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 40, Application US/10307276B Publication No. US20040101904A1 GENERAL INFORMATION:
    Query Match
                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/864,408A
CURRENT FILING DATE: 2001-05-24
PRIOR APPLICATION NUMBER: 60/206,690
PRIOR FILING DATE: 2000-05-24
                                                         TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                          NUMBER OF SEQ ID NOS: 9068
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/307,276B

FILING DATE: 27-NOV-2002

CLASSIFICATION: «Unknown»

ATTORNEY/AGENT INFORMATION:

NAME: Oldenkamp, David J.

REGISTRATION NUMBER: 29,421

REGISTRATION NUMBER: 0180.0038

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: polypeptide
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS: LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Shapiro & Dupont LLP
STREET: 233 Wilshire Boulevard, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Delivery Of Pharmaceutical Agents
Via The Human Insulin Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: William M. Pardridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 2000
SOFTWARE: MS Word
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (310) 319-5411
TELEFAX: (310) 319-5401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 90067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Santa Monica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
  100.0%; Score 31;
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Pred. No. 1.3e+06;
); Mismatches 0;
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  DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 7;
Length 97;
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CURRENT APPLICATION NUMBER: US/10/010,729
CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 09/730,473
PRIOR FILING DATE: 2000-12-05
PRIOR PELICATION NUMBER: 09/580,787
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 09/322,862
PRIOR APPLICATION NUMBER: 09/322,862
PRIOR PELING DATE: 199-05-28
PRIOR PELICATION NUMBER: 08/79,784
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR FILING DATE: 1996-08-08
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR FILING DATE: 1994-04-29
PRIOR FILING DATE: 1994-04-29
NUMBER OF SEO ID NOS: 80
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 45
LENGTH: 108
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/09/924,099
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/338,511
PRIOR FILING DATE: EARLIER FILING DATE: 1999-06-23
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: JP 289,044/98
PRIOR APPLICATION NUMBER: EARLIER FILING DATE: 1998-10-12
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: JP 365,023/98
PRIOR PRIOR EILING DATE: EARLIER FILING DATE: 1998-12-22
PRIOR PRIOR EARLIER FILING DATE: 1998-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Human IgM Antibodies and Diagnostic and TITLE OF INVENTION: Therapeutic Uses Thereof Particularly in the Central Nervous TITLE OF INVENTION: System FILE REFERENCE: 1199-1-005CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Rodriguez, Moses APPLICANT: Miller, David J APPLICANT: Pease, Larry R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 33
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TYPE: PRT
ORGANISM: Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 100.0%; Score 31; DB Similarity 100.0%; Pred. No. 34; 7; Conservative 0; Mismatches
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KURIMOTO, Masashi
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; ORGANISM: Mus musculus
US-10-010-729-45
                                                                                                                   US-10-307-276B-6
                                                                                                                                       RESULT 7
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US-10-307-276B-4
                                                            Sequence 6, Application US/10307276B Publication No. US20040101904A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/10307276B Publication No. US20040101904A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                      Query Match
Best Local 8
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                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acids
TYPE: amino acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 2000
SOFTWARE: MS WOR'D
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Oldenkamp, David J.
REGISTRATION NUMBER: 29,421
REPERENCE/DOCKET NUMBER: 0180.0038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (310) 319-5411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/10/307,276B
FILING DATE: 27-Nov-2002
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ruben J. Boado
TITLE OF INVENTION: Delivery Of Pharmaceutical Agents
Via The Human Insulin Receptor
Ruben J. Boado
TITLE OF INVENTION: Delivery Of Pharmaceutical Agents
                                       APPLICANT: William M. Pardridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 90067
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESSE: Shapiro & Dupont LLP
STREET: 233 Wilshire Boulevard, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: William M. Pardridge
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                                                                                                                                                                                                                                                                     Similarity 7; Conserv
                                                                                                                                                                                            ATSSLDS 56
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STATE: CA
COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (310)
                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                     100.0%; Score 31; DE
100.0%; Pred. No. 34;
vative 0; Mismatches
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Pred. No. 34;
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                                                                                                                                                                                                                                                                                                        DB 16; Length 108;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 267, Application US/10803622
Publication No. US20040157214A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
             APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OP INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REPERSUE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 902845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
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                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/10/307,276B
FILING DATE: 27-Nov-2002
CLASSIFICATION: <UNKnown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 90067
COMPUTER READABLE FORM:
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Similarity 100.0%;
7; Conservative 0
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REGISTRATION NUMBER: 29,421
REFERENCE/DOCKET NUMBER: 0180.0038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: Windows 2000 SOFTWARE: MS Word
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy Disk
                                                                                                                                                                                                          Marks, James
Clackson, Timothy
Chiswell, David
                                                                                                                                                                                                                                                                   Jackson, Ronald
Holliger, Kasper
                                                                                                                                                                                                                                                                                                        Hoogenboom, Hendricus
Griffiths, Andrew
                                                                                                                                                                                                                                                                                                                                           Johnson, Kevin
                                                                                                                                                                                                                                                                                                                                                                 Pope, Anthony
                                                                                                                                                                                                                                                                                                                                                                                   McCafferty, John
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Pred. No. 34;
0; Mismatches
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NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESSE: Shapiro & Dupont LLP
STREET: 233 Wilshire Boulevard, Suite 700
CITY: Santa Monica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Via The Human Insulin Receptor
                                 16; Length 108;
   Indels
   0
Gaps
   0
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APPLICANT: Winter, Gregory
APPLICANT: Winter, Timothy
ITITE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERICE; 11803-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR RILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR RILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR PILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1990-13-06
PRIOR FILING DATE: 1990-03-06
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PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR APPLICATION NUMBER: PCT/GB91/01134
PRIOR APPLICATION NUMBER: DOT/971,857
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 267, A
Publication No.
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SEQ ID NO 267
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Cambridge Antibody Technology Limited
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: light chain from clone M1F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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les 7; Conserv
                 APPLICATION NUMBER: US 07/971,857 FILING DATE: 1993-01-08
                                                           FILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 1990-10-19
APPLICATION NUMBER: GB 9024503.6
FILING DATE: 1990-11-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: GB 9104744.9
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Marks, James
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Griffiths, Andrew
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DS20040157215A1
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  NUMBER:
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08/484,893
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; MOLECULE TYPE: peptide ; SEQUENCE DESCRIPTION: 0US-09-943-906-74
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Best Local Similarity
"hes 7; Conserve
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   Matches
                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 74, Application U Patent No. US20020150571A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 272
SOFTWARE: PatentIn version 3.1
SEQ ID NO 267
LENGTH: 108
TYPE: PRT
                                                                                                                                                                        TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 74:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: light chain
 Local Similarity
nes 7; Conserv
                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Bozicevic, Karl
REGISTRATION NUMBER: 28,807
                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/943,906
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road
CITY: Menlo Park
                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Williamson, R. Anthony
Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Prusiner, Stanley B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50
                                                                                                                          STRANDEDNESS: single
                                                                                                                                          TYPE: amino acid
                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version 2.0
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                                                                                                                                                           LENGTH: 109 amino acids
                                                                                                                                                                                                                                          TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: CA
                                                                                                                                                                                                                           TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94025
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 Conservative
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                100.0%; Score 31; DB 100.0%; Pred. No. 34;
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                                                                              SEQ ID NO:
 0
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Pred. No. 34;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 from clone M1F
Mismatches
                                                                                                                                                                                                                                                                          06510/059001
                                                                                74:
                              DB 9; Length 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 16;
 0
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 Indels
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 0,
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Gaps
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ATSSLDS

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50 ATSSLDS 56

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APPLICANT: FURBINET, Stanley
APPLICANT: Safar, Jiri
APPLICANT: Williamson, Anthony
APPLICANT: Williamson, Anthony
APPLICANT: Williamson, Anthony
ITITLE OF INVENTION: Antibodies Specific for Ungulate PrP
FILE REFERENCE: UCAL-194
CURRENT APPLICATION NUMBER: US/10/355,780
CURRENT FILING DATE: 2003-01-30
PRIOR APPLICATION NUMBER: US/09/627,218B
PRIOR APPLICATION NUMBER: US/09/627,218B
PRIOR APPLICATION NUMBER: US/09/627,218B
PRIOR APPLICATION NUMBER: US/09/627,218B
PRIOR FILING DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 11
SOPTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                             ; OTHER INFORMATION: synthesized peptide
US-10-355-780-1
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US-10-355-780-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-435-602-74
; Sequence 74, Application US/10435602
; Publication No. US20030228303A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 09/943,906
PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/550,374
PRIOR FILING DATE: 2000-04-13
PRIOR PRIOR DATE: 2000-04-13
PRIOR PILING DATE: 1998-03-06
PRIOR PILING DATE: 1998-03-06
PRIOR PILING DATE: 1996-09-13
PRIOR PILING DATE: 1996-09-13
PRIOR PILING DATE: 1996-09-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/10355780 Publication No. US20030143224A1 GENERAL INFORMATION:
                                                                                       Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOPTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 74
                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Prusiner, Stanley B.
APPLICANT: Williamson, R. Anthony
APPLICANT: Williamson, R. Anthony
APPLICANT: Burton, Dennis R.
TITLE OF INVENTION: Antibodies Specific for Native PrPSc
PILE REFERENCE: UCAL059CON3
CURRENT APPLICATION NUMBER: US/10/435,602
CURRENT FILING DATE: 2003-05-09
CURRENT FILING DATE: 2003-05-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1995-09-14
                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                                                                                                                                                                                                       ENGTH: 112
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nes 7; Conserv
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1 ATSSLDS 7
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                                                                     Similarity 7; Conser
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                                                                100.0%; Score 31; DB 14; llarity 100.0%; Pred. No. 35; Conservative 0; Mismatches 0;
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Pred. No. 34;
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                                                                                                            Length 112;
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; HYPOTHETICAL: NO
; FRAGMENT TYPE; N
; ORIGINAL SOURCE;
; ORGANISN; MUS
US-08-779-784-35
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RESULT 14
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US-08-779-784-35
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Publication No.
                                                                                                                     Matches
                                                                                                                                 Best Local
                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 201-343-1684
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Jackson Esq., David A. REGISTRATION NUMBER: 26,742 REFERENCE/DOCKET NUMBER: 119: TELECOMMUNICATION INFORMATION: TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 08-AUG-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/01
FILING DATE: 07-JAN-1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
APPLICATION NUMBER: US 01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Miller, David J.
APPLICANT: ASSAURA, KUMINIKO
TITLE OF INVENTION: PROMOTION OF CENTRAL NERVOUS SYSTEM
TITLE OF INVENTION: REMYELINATION USING MONOCLONAL AUTOANTIBODIES
                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 29-APR-1994
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Hackensack
STATE: New Jersey
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                                                 72 ATSSLDS 78
                                                                                1 ATSSLDS 7
                                                                                                                   Similarity 100.0%; 7; Conservative (
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                                                                                                                 Score 31; DB
Pred. No. 41;
0; Mismatches
                                                                                                                                                   DB 8; Length 130;
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                                                                                                                 Gaps
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Sequence 71, Application US/1
Publication No. US2003018587
GENERAL INFORMATION:
APPLICANT: Rodriguez, Moses
APPLICANT: Miller, David J.

Application US/10010729 10. US20030185827A1

US-10-010-729-71

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GENERAL INFORMATION:

APPLICANT: Thorpe, Philip E.

APPLICANT: Soares, M. Melina

APPLICANT: Soares, M. Melina

APPLICANT: Soares, M. Melina

ITILE OF INVENTION: Methods for Treating Viral Infections Using Antibodies to

TITLE OF INVENTION: Methods for Treating Viral Infections Using Antibodies to

TITLE OF INVENTION: Methods for Treating Viral Infections Using Antibodies to

TITLE OF INVENTION: Methods for Treating Viral Infections Using Antibodies to

TITLE OF INVENTION: METHODS WITH INFECTION UNMERS: US/10/642,120

CURRENT FILING DATE: 2003-08-15

PRIOR APPLICATION NUMBER: US/10/642,120

CURRENT FILING DATE: 2003-07-15

PRIOR APPLICATION NUMBER: 60/396,263

PRIOR FILING DATE: 2002-07-15

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin version 3.1

SEQ ID NO 4

LENGTH: 144

TYPE: PRT

ORGANISM: Mus musculus
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TITLE OF INVENTION: Human 1gm Antibodies and Diagnostic and
TITLE OF INVENTION: Therapputic Uses Thereof Particularly in the Central Nervous
TITLE OF INVENTION: System
FILE REFERENCE: 1199-1-005CIP2
CURRENT PRICKING DATE: 2000-12-05
PRIOR APPLICATION NUMBER: 09/730,473
PRIOR FILING DATE: 2000-12-05
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR APPLICATION NUMBER: 09/582,862
PRIOR FILING DATE: 1999-05-28
PRIOR FILING DATE: 1999-05-28
PRIOR FILING DATE: 1990-05-28
PRIOR FILING DATE: 1990-01-07
PRIOR APPLICATION NUMBER: 08/739,784
PRIOR FILING DATE: 1997-01-07
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR FILING DATE: 1997-01-07
PRIOR APPLICATION NUMBER: 08/236,520
PRIOR APPLICATION NUMBER: 08/236,520
PRIOR APPLICATION NUMBER: 08/236,520
PRIOR APPLICATIO
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US-10-642-120-4
; Sequence 4, Application US/10642120
; Publication No. US20040131610A1
; Publication No. US20040131610A1
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Search completed: April 4, 2005, 16:42:45 Job time: 11.3323 secs
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Maximum Match 100%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2005 Compugen Ltd.
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C;Accession: Fu0260

R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A.

J. Exp. Med. 171, 265-297, 1990

A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic n
A;Reference number: PL0231; MUID:90111618; PMID:2104919

A;Accession: PL0260

A;Molecule type: mRNA
A;Residues: 1-106 <SHL>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;14-23/Region: framework 1
F;16-90/Domain: immunoglobulin homology <TMM>
F;24-34/Region: complementarity-determining 1
F;35-49/Region: complementarity-determining 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 2
PL0260
If kappa chain V region (anti-DNA, DP7VK) - mouse (fragment)
If kappa chain V region (anti-DNA, DP7VK) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
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If kappa chain V region (anti-DNA, DP15VK) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 09-May-1997
C;Accession: PLO261
R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A. J. Exp. Med. 171, 265-297, 1990
A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic n A;Reference number: PLO231; MUID:90111618; PMID:2104919
A;Accession: PLO261
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A;Residues: 1-88 <SHL>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotettamer; immunoglobulin
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F;80-88/Region: framework 4
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F;32-38/Region: complementarity-determining
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F;57-88/Region:
F;89-97/Region:
F;98-106/Region:
A;Cross-references: GB:V00804; GB:J00566; NID:g52127; PIDN:CAA24186.1; FA;Note: the sequences were determined from the differentiated gene MOPC R;Burstein, Y.; Schechter, I.
Proc. Natl. Acad. Sci. U.S.A. 74, 716-720, 1977
A;Title: Amino acid sequence of the NH-2-terminal extra piece segments cal; Reference number: A93815; MUID:77148916; PMID:403522
A;Accession: A93815
A;Molecule type: protein
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A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal A;Reference number: PL0231; MUID:90111618; PMID:2104919

A;Accession: PL0259
                                                                                                                                                                                                                                                                         R;Seidman, J.G.; Max, B.B.; Leder, P.
Nature 280, 370-375, 1979
A;Title: A kappa-immunoglobulin gene is formed by site-specific A;Reference number: A93211; MUID:79221900; PMID:111146
A;Accession: A93211
A;Molecule type: DNA
A;Residues: 1-130 <PC41>
                                                                                                                                                                                                                                                                                                                                                                                                                Ig kappa chain precursor V region (MOPC 41) - mouse N;Contains: Ig kappa chain precursor V region VK41 C;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 24-Apr-1984 #sequence revision 24-Apr-1984 #text_change 09-Jul-2004 C;Date: 24-Apr.1984 #sequence revision 24-Apr-1984 #text_change 09-Jul-2004 C;Accession: A93211; B93211; A93815; A94239; A01922; A01923 R;Seidman, J.G.; Max, E.E.; Leder, P.
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A; Residues: 1-106 < SHL>
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A; Residues: 1-117 < VK41>
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;Species: Mus musculus (house mouse)
;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
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hypothetical protein RC0301 [imported] - Rickettsia conorii (strain Malish 7) C; Species: Rickettsia conorii (c; Species: Rickettsia conorii (c; Species: Rickettsia conorii (strain Malish 7) C; Species: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004 C; Accession: E97737 R.Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P.E.; Barbe, V.; Samson, Science 293, 2093-2098, 2001 R.J.Title: Mechanisms of Evolution in Rickettsia conorii and Rickettsia prowazekii. A; Reference number: A97700; MUID:21442074; PMID:11557893 A; Accession: E97737 A; Reference E97737 A; Status: preliminary A; Status: preliminary
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Science 155, 465-467, 1967
A;Title: Mechanism of antibody synthesis: 6
A;Reference number: A94239; MUID:67056897;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:Q92IW9; GB:AE006914; PIDN:AAL02839.1; PID:g15619360; GSPDB:GPC;Genetics:
A;Gene: RC0301
C;Superfamily: multidrug resistance protein; ATP-binding cassette homology
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A;Residues: 23-49,'B',51-53,'LSB',57-58,'ZZ',61-62,'BZ',65-76,'B',78-108,110-130 <GRA:
A;Experimental source: Bence Jones protein MOPC 41
                        R;DelVecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan, Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucell A;Reference number: AD3252; PMID:11756688
A;Accession: AI3469
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A; Status: preliminary
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; PMID:4162931
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C/Suportamily: ... C;Keywords: ATP
C;Keywords: ATP
F;323-517/Domain: ATP-binding cassette homology <ABC>
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A; Residues: 1-555 < OBE>
A; Cross-references: UNIPROT: P53877;
A; Cross-references: Btrain S288C
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A;Accession: F71732
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                                                                                                                                                                                                                                                                             submitted to the Protein Sequence A; Reference number: S63122 A; Accession: S63137
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                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein YNL182c - yeast (Saccharomyces cerevisiae)
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A; Residues: 1-548 < AND>
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C;Spocies: Rickettsia prowazekii
C;Date: 21-Nov-1998 #sequence_revision 21-Nov-1998 #text_change 16-Aug-20
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Superfamily: ATP-b
                                                                                                                                                        Cross-references:
                                                                                                                                                                          Gene: MIPS:YNL182c
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                                                                                      Matches
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R;Kunst, F; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berterc C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Choj A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E. Nature 390, 249-256, 1997

A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallerc iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F.; Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, A.; Ahthors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maucel, Y., M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle, Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scandon, A.; Aputhors: Schleich, S.; Schroetter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Serror, akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama, T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K. A; Authors: Yoshikawa, H.; Danchin, A.
A;Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis. A; Rocession: D70031
                                                                                                                                                                                                                                                                                                                                                                                                                        R;Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P.E.; Barbe, V.; Samson, Science 293, 2093-2098, 2001
S,Title: Mechanisms of Evolution in Rickettsia conorti and Rickettsia prowazekii. A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: F97735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein abcT3 [imported] - Rickettsia conorii (strain Malish 7 C;Species: Rickettsia conorii C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 16-Aug-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Superfamily: Escherichia coli ABC transporter mdlA; ATP-binding C;Keywords: ATP; nucleotide binding; P-loop F;357-552/Domain: ATP-binding cassette homology <ABC> F;374-381/Region: nucleotide-binding motif A (P-loop)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABC transporter (ATP-binding protein) homolog yvcC - Bacillus subtilis C;Species: Bacillus subtilis C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004 C;Accession: D70031
                                                                                                                                                                                                A; Gene: abcT3
C; Superfamily:
                                                                                                                                                                                                                                                                                            A;Molecule type: DNA
A;Residues: 1-589 <KUR>
A;Cross-references: UNIPROT:Q92IY4; GB:AE006914;
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A;Cross-references: UNIPROT:006967; GB:Z99121; GB:AL009126; NID:g2635827; PIDN:CAB15487.1
A;Experimental source: strain 168
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Pred. No. 15;
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ATSSLDS 517

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Proc. Nat
A;Title:
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A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacterium tum A;Reference number: A97359; MUID:21608551; PMID:11743194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Agrobacterium tumefaciens
C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 16-Aug-2004
C;Accession: H98149
                                       R; Finan,
                                                       msbA-like saccharide exporting ABC transporter protein, consisti C;Species: Sinornizobium meliloti C;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change C;Accession: H95974
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A;Molecule type: DNA
A;Residues: 1-600 <KUR>
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A; Residues: 1-600 < KUR>
A; Cross-references: UNIPROT: Q8U6S8;
A; Experimental source: strain C58 (D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Pester, E.W.
B;Title: The Genome of the Natural Genetic Engineer Agrobacterium A;Reference number: AB2577; MUID:21608550; PMID:11743193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, t, karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Agrobacterium tumefaciens
C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change
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Best Local S
Matches 7
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in, T.M.; Weidner, S.; Wong, In, T.M.; Weidner, S.; Wong, Natl. Acad. Sci. U.S.A. 98,
                                                                                                                                                                                                                                                                                                                                                                    position:
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 K.; Buhrmester, J.; Chain,
9889-9894, 2001
the 1,683-kb pSymB megaplas
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(Dupont)
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                                     P.; Vorholter,
                                                                           16-Aug-2004
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RESULT 14
S60182
              R;Rubio, J.P.; Cowman, A.F.
Exp. Parasitol. 79, 137-147, 1994
Exp. Parasitolum falciparum: the pfmdr2
A;Reference number: Z18924; MUID:94333528;
                                                                                                              multidrug resistance protein 2 - malaria parasite C;Species: Plasmodium falciparum C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 C;Accession: T18376
A; Reference number: A; Accession: T18376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Superfamily: ATP-binding cassette homology C;Keywords: ATP; nuclectide binding; P-loop F;371-565/Domain: ATP-binding cassette homology <ABC>F;388-395/Region: nuclectide-binding motif A (P-loop)
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A;Cross-references: UNIPROT:Q52924; EMBL:Z50189; NID:g1143532; PIDN:CAA90568.1; PID:e1914
A;Note: it is uncertain whether Met-1 (ATG), Val-29 (GTG) or Met-74 (ATG) is the initiate
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A;Title: Extension of the Rhizobium meliloti succinoglycan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATP-binding transport protein exsA - Rhizobium meliloti
C;Species: Rhizobium meliloti
C;Date: 15-Feb-1996 #sequence_revision 01-Mar-1996 #text_change 16-Aug-2004
C;Accession: S60182
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A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, hebault, P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, A;Title: The composite genome of the legume symbiont Sinorhizobium meliloti A;Reference number: A96039; MUID:21368234; PMID:11474104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Residues: 1-604 < XUR>
A;Cross-references: UNIPROT:Q52924; GB:AL591985; PIDN:CAC49464.1; PID:g15140950; GSPDB:GPA;Cross-references: UNIPROT:Q52924; GB:AL591985; PIDN:CAC49464.1; PID:g15140950; GSPDB:GPA;Experimental source: strain 1021, megaplasmid pSymB
R;Galibert, F.; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barloy-Hubler, Pela, D.; Chain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fisher, R.F.; L.; Hyman, R.W.; Jones, T.
L.; Hyman, R.W.; Jones, T.
Science 293, 668-672, 2001
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A;Accession: H95974
A;Status: preliminary
A;Molecule type: DNA
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A.Secular Proliminary, translated from GB/BNEL/DDB3
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A.PROMOMOR O'PPOON MAND.
A.COMORT GENERAL MAND.
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ALIGNMENTS

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RESULT 2

KV5G MOUSE

KV5G MOUSE

ID

KV5G MOUSE

STANDARD;

PRT; 130 AA.

AC

P01639; P01640;

PT

21-UUL-1986 (Rel. 01, Created)

PT

21-UUL-1986 (Rel. 01, Last sequence update)

PT

05-JUL-2004 (Rel. 44, Last annotation update)

DE

IG

Kappa chain V-V region MOPC 41 precursor.

OS

Mus musculus (Mouse).

OC

Eukaryota; Metazoa; Chordata; Craniata; Vertebra

OC

Mammalia; Eutheria; Rodentia; Sciurognathi; Muri

OX

NCBI TaxID=10090;

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EMBL; AABW0100001; EAA25646.1; -.

GO; GO:0016202; C:membrane; IEA.

GO; GO:001622; F:ATP binding; IEA.

GO; GO:000524; F:ATP binding; IEA.

GO; GO:0042626; F:ATPase activity, coupled to trar

GO; GO:0006810; P:transport; IEA.

InterPro; IPR003439; ABC_transporter.

PROSITE; PS00211; ABC_TRANSPORTER_1; UNKNOWN_1.
   Ig kappa chain V-V region MOPC 41 precursor.
Mus musculus (Mouse).
Eukaria, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI TaxID=10090;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Malek J.A., Eremeeva M.E., Dasch G.A.;
Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the ABC transporter family.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales; Rickettsiaceae; Rickettsieae; Rickettsia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Multidrug resistance ABC transporter ATP-binding protein. Name=rsib_orf.406;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 31; DB 2; Length 57; Pred. No. 6.8; Mismatches 0; Indels
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RESULT
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Best Local Similarity
"hehes 7; Conservi
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MEDLINE=79221900; PubMed=111146;
MAY R.E., Leder P.;
                                                                                                                                                                                                                                                                      Q92IW9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
DISULFID
VARIANT
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Burstein Y., Schechter I.; Pamino acid sequence of the NH2-terminal extra piece segments of precursors of mouse immunoglobulin lambdal-type and kappa-type 1?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seidman J.G., Max E.E., Leder P.; "A kappa-immunoglobulin gene is for without further somatic mutation.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOMAIN
DOMAIN
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Bence-Jones protein; Direct p:
Immunoglobulin V region; Sign:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- MISCELLANEOUS: This precursor was synthesized system directed by mRNA isolated from myeloma-!- MISCELLANEOUS: This is a Bence-Jones protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chains."
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                                                                                                                                                                                    01-DEC-2001 (TYENHELTE1. 19, Created)
01-DEC-2001 (TYENHELTE1. 19, Last sequence update)
01-DEC-2003 (TYENHELTE1. 25, Last annotation update)
Multidrug resistance ABC transporter ATP-binding pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=67056897;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                              Rickettsia conorii.
                                                                                                                                                                   OrderedLocusNames=RC0301;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DOMAIN
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  MEDLINE=21442074; PubMed=11557893; DOI=10.1126/science.1061471;
                       SEQUENCE FROM N.A. STRAIN=Malish 7;
                                                                                                           Rickettsiaceae; Rickettsieae;
                                                                                                                        Bacteria; Proteobacteria;
                                                                             VCBI_TaxID=781;
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gene is formed
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Pred. No. 18;
); Mismatches
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Missing (in
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                                                                                                           Rickettsia.
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                                                                                                                              Rickettsiales;
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GO; GO:0016024; F:AIP binding; IEA.
GO; GO:00042626; F:AIPsea activity, coupled to
GO; GO:00042626; F:AIPsea activity, coupled to
GO; GO:0006810; P:transport; IEA.
GO; GO:0006810; P:transporter.
InterPro; IPR003439; ABC_transporter.
Pfam; PF000005; ABC_transporter; 1.
ProDom; PD000006; ABC_transporter; 1.
PROSITE; PS00211; ABC_TRANSPORTER_1; 1.
ATP-binding; Complete protecome.
SEQUENCE 131 AA; 14753 MW; C8484BCCA61287
                   Query Match
Best Local Similarity
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Samson D.,
Raoult D.;
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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

STRAIN=16M / ATCC 23456 / Biotype 1;

STRAIN=20020109; PubMed=11756688; DOI=10.1073/pnas.221575398;

MEDLINE=20020109; PubMed=11756688; DOI=10.1073/pnas.221575398;

Los DelVecchio V.G., Kapatral V., Redkar R.J., Patra G., Mujer C., Los DelVecchio V.G., Kapatral V., Edakar R.J., Lykidis A., Reznik G., Ivanova N., Anderson I., Bhattacharyya A., Lykidis A., Reznik G., Ivanova N., Anderson I., Bhattacharyya A., Lykidis A., Reznik G., Ivanova N., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M., Goltsmar M., Ballaghan D., Letesson J.-J., Jablonski L., Larsen N., Ballaghan D., Letesson J.-J., Jablonski L., Larsen N., Bernal A., Mazur M., Ballaghan D., Letesson J.-J., Larsen N., Ballaghan D., Letesson J.-J., Mazur M., Ballaghan D., Letesson M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "The genome sequence of the facultative intrac
Brucella melitensis.";
Proc. Natl. Acad. Sci. U.S.A. 99:443-448(2002)
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                                                                               PROSITE; PS00211; ABC_TRANSPORTER_1; PROSITE; PS50893; ABC_TRANSPORTER_2; ATP-binding; Complete proteome. SEQUENCE 320 AA; 35768 MW; 95DBAB
                                                                                                                                                                                                     GO; GO:0000166; F:nucleotide binding; GO; GO:0006810; P:transport; IEA. Pfam; PF00005; ABC tran; 1. Probom; PD00006; ABC transporter; 1. SMART; SM00382; AAA; 1.
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HSSP; Q9KQW9; 1PF4.
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Haselkorn R.,
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GO:0005524; F:ATP binding; IEA
GO:0042626; F:ATPase activity;
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Roux V.,
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R., Kyrpides N.C., C
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Last annotation update)
proTEIN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           overbeek R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 31;
Pred. No.
   Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                              IEA.
                                                                                            95DBAB38430A2C01 CRC64;
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Mismatches '
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                                   Length
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                                      320;
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1., Goltsman
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IEA.

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RESULT

ID 11009

AC 061

AC 060

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ID Q81E
AC Q81E
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DT 01-J
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RA Ivan
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Best Local
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Q61009;
05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
ABC transporter.
                                                                                                                        PROSITE; PS50929; ABC_TM1F; 1.
PROSITE; PS00211; ABC_TRANSPORTER_2; 1.
PROSITE; PS50893; ABC_TRANSPORTER_2; 1.
ATP-binding.
SEQUENCE 370 AA; 40954 MW; 9C668329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pfam; PF00664; ABC_membrane; 1.
Pfam; PF00005; ABC_tran; 1.
ProDom; PD000006; ABC_transporter; 1.
SMART; SM00382; AAA; 1.
   SEQUENCE FROM N.A.
MEDLINE=22608415; PubMed=12721630; DOI=10.1038/nature01582; Ivanova N., Sorokin A., Anderson I., Galleron N., Candelon
                                                                                       Bacillus cereus (strain ATCC 14579
Bacteria; Firmicutes; Bacillales;
                                                                                                                      protein.
OrderedLocusNames=BC1955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR003593; AAA ATPase.
InterPro; IPR011527; ABC membrane 1.
InterPro; IPR001140; ABC_TM_transpt.
InterPro; IPR003439; ABC_transporter.
                                                                     NCBI_TaxID=226900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OrderedLocusNames=BAS1815;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN-Sterne;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacillus anthracis.
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                                                                                                                                                                                                                                                                                                                                      ATSSLDS 296
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                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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Last annotation updat
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                                                                                         14579 / DSM 31).
mles; Bacillaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9C6683290C2C7EC3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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c., Okinaka J
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Q9ZDV5
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R PROSITE; PS50929; ABC_TRANSPORTER_1; 1.

OR PROSITE; PS50893; ABC_TRANSPORTER_2; 1.

PROSITE; PS50893; ABC_TRANSPORTER_2; 1.

PROSITE; PS50893; ABC_TRANSPORTER_2; 1.

PROSITE; PS50893; ABC_TRANSPORTER_2; 1.
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Best Local S
Matches 7
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Q9ZDV5;
01-MAY-1999
                                                                                                                                                      -!- SIMILARITY: Belongs to EMBL; AJ235270; CAA14677.1; PIR; F71732; F71732.
                                                                                                                                                                                                                                           MEDLINE=99039499; PubMed=9823893; DOI=10.1038/24094; Andersson S.G.E., Zomorodipour A., Andersson J.O., Sicheritz-Ponten T., Alsmark U.C.M., Podowski R.M., Naeslund Eriksson A.-S., Winkler H.H., Kurland C.G.; "The genome sequence of Rickettsia prowazekii and the origin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
01-OCT-2003 (TrEMBLrel. 25,
ABC TRANSPORTER ATP-BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88888
                                                                                                                                                                                                              Nature 396:133-140(1998).
                                                                                                                                                                                                                              "The genome sequence mitochondria.";
                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. STRAIN=Madrid E;
                                                                                                                                                                                                                                                                                                                                                                                                       Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiaceae; Rickettsieae; Rickettsia.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Rickettsia prowazekii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     OrderedLocusNames=RP214;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR003593; AAA ATPase.
InterPro; IPR011527; ABC membrane 1.
InterPro; IPR001140; ABC TM transpt.
InterPro; IPR003439; ABC transporter
Pfam; PF00664; ABC membrane; 1.
Pfam; PF00005; ABC tran; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -i- SIMILARITY: Belongs to the EMBL; AE017004; AAP08926.1; -HSSP; Q9CHL8; 1MV5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kapatral V., Bhattacharyya A., Reznik G., Mikhailova N., Lapidu
Chu L., Mazur M., Goltsman E., Larsen N., D'Souza M., Walunas T
Grechkin Y., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,
InterPro; IPR003593; AAA_ATPase.
InterPro; IPR011527; ABC_membrane_1.
InterPro; IPR001140; ABC_TM_transpt.
                                                                                                                                        HSSP; P08716; 1MT0
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Nature 423:87-91(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Grechkin Y., Pusch G., Haselkorn Overbeek R., Kyrpides N.C.;
                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=782;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Genome sequence of Bacillus cereus and
                                                    GO:0016021; C:integral to membrane; IBA. GO:0005524; F:ATP binding; IBA. GO:0042626; F:ATPase activity, coupled t GO:0000166; F:nucleotide binding; IEA. GO:0006810; P:transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GO:0016021; C:integral to membrane;
GO:0005524; F:ATP binding; IEA.
GO:0042626; F:ATPase activity, coup.
GO:0000166; F:nuclectide binding; II
GO:0006810; P:transport; IEA.
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Last annotation update)
PROTEIN (AbcT3).
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Pred. No.
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                                                                                         coupled to transmembrane
                                                                                                                                                                                            transporter
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                                                                                                                                                                                            family
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IEA.

ProDom; PD000006; ABO SMART; SM00382; AAA;

Pfam; PF00664; ABC_membrane; Pfam; PF00005; ABC_tran; 1. Pr0D00m; PD000006; ABC transpc SMART; SM00382; AAA; 1.

transporter;

IPR003439; ABC_transporter.

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YNS2
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ID YNS3
ID YNS3
ID YNS2
AC P538
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Best Local S
Matches 7
                                                                                                                                                                                                                                    InterPro: IPRO01680; WD40.

InterPro: IPRO11046; WD40_like.

Pfam; PP00400; WD40; 2.

SMART; SM00320; WD40; 3.

PROSITE; PS00678; WD REPEATS_1; FALSE_NEG.
PROSITE; PS0062; WD REPEATS_2; 1.

PROSITE; PS50024; WD_REPEATS_REGION; 1.

Hypothetical protein; Repeat; WD repeat.

REPEAT 137 176 WD 2.

REPEAT 137 176 WD 2.
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YEAST
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STANDARD;
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555 AA
P53877;
01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation updat
Hypothetical 61.8 kDa Trp-Asp repeats conta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE; PS50929; ABC_TM1F; 1.
PROSITE; PS00211; ABC_TRANSPORTER_1; 1.
PROSITE; PS50893; ABC_TRANSPORTER_2; 1.
ATP-Dinding; Complete proteome.
SEQUENCE 548 AA; 62589 MW; B4ED4E7F9D53EDC9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.

Obermaier B., Piravandi E., Rinke M., Domdey H.;
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OrderedLocusNames=YNL182C; ORFNames=N1636; Saccharomyces cerevisiae (Baker's yeast). Saccharomyces cerevisiae (Baker's yeast). Bukaryota; Fungi, Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; Z71458; CAA96075.1; PIR; S63137; S63137.
                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SGD; S000005126; IPI3.
GO; GO:0005634; C:nucleus; IDA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=4932;
                                                                                                                                                                                                                      REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GermOnline; 143188;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IntAct; P53877;

    -!- SIMILARITY: Contains 4 WD repeats.

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                                                                                                                                                                 555 AA;
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100.0%; Pred. No. 92;
cive 0; Mismatches
                                                       0;
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4. 3. 2. 1.
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Pred. No.
                                                                                                                                                              8EED6854DF9405A6 CRC64;
                                                       Mismatches
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containing
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                                                                                                         Length 555;
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RESULT 11

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Best Local S
Matches 7
                                                                                  J. VIrol. 74:3464-3469 (2000).
J. VIrol. 74:3464-3469 (2000).
EMBL; AJ270057; CAB85589.1; -.
GO; GO:0005198; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecule
InterPro; IPR003514; Caspid_F.
Pfam; PF02305; Phage_F; 1.
SEQUENCE 565 AA; 63538 MW; 670485C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q6KF06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Garner S.A., Everson J.S., Lambden P.R., Fane Submitted (MAR-2003) to the EMBL/GenBank/DDBJ EMBL, AJ550635, CAD79477.1; ... GO; GO:0019028; C:viral capsid; IEA. GO; GO:0005198; F:structural molecule activity InterPro; IPR003514; Caspid_F. PF02305; Phage_F: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Structural protein.
Chlamydia phage 3.
Viruses; ssDNA viruses; Microviridae; Chlamydiamicrovirus
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01-OCT-2000 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                               Viruses; ssDNA viruses;
NCBI_TaxID=105154;
                                                                                                                                                                                                                                                                                                                                       дэмвие;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=225067;
                                                                                                                                                              psittaci."
                                                                                                                                                                                 DOI-10.1128/JVI.74.8.3464-3469.2000;
Liu B., Everson J.S., Fane B., Giannikopoulou P.,
Lambden P.R., Clarke I.N.;
                                                                                                                                                                                                                    MEDLINE=20193783; PubMed=10729119;
                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                       Chlamydia phage 2.
                                                                                                                                                                                                                                                                                              Structural protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                          "Molecular characterisation of a bacteriophage (Chp2) from Chlamydia
                                                                                                                                                                                                                                                                                                                                                                          10
                                                                                                                                                                                                                                                                                                                                                                                                          497
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                   ATSSLDS 7
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 ATSSLDS 503
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                                           Conservative
                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 31; 100.0%; Pred. No.
                                                    100.0%;
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15,
26,
                                                                                                                                                                                                                                                                Microviridae;
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Last annotation update)
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Last annotation update)
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                                                                                                                     molecule activity;
                                                    Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                     PRT;
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                                                                                     670485CDC2A94932 CRC64;
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                                            Mismatches
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RESULT 12
0739UB
Q739UB
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AC Q739UB
D7 05-J1
D7 0
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PubMed-14960714; DOI=10.1093/nar/gkh258;
Rasko D.A., Ravel J., Oekstad O.A., Helgason E., Cer R.Z., Jiang L.,
Shores K.A., Fouts D.E., Tourasse N.J., Angiuoli S.V., Kolonay J.F.,
Nelson W.C., Kolstoe A.-B., Fraser C.M., Read T.D.;
"The genome sequence of Bacillus cereus ATCC 1997 reveals metabolic
adaptations and a large plasmid related to Bacillus anthracis pXOI.",
Nucleic Acids Res. 32:977-988(2004).
--- SIMILARITY: Belongs to the ABC transporter family.
EMBL; AE017270; AAS40963.1; -.
EMBL; AE017270; AAS40963.1; -.
GO; GO:0016021; C:integral to membrane; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q739U8;
Q739U8;
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ATP-binding.
SEQUENCE 586 AA; 64902 MW; BCE45D1F31C7CE7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation updat
ABC transporter, ATP-binding/permease protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brettin T.S., Bruce D., Challacom
Hitchcock P., Jackson P., Keim P.
Richardson P., Rubin B., Tice H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacillus cereus (strain ATCC 10987).
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OrderedLocusNames=BCE2040;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacillus cereus ZK.
Bacteria, Pirmicutes; Bacillales; Bacillaceae, Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN-ZK;
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Pred. No. 1e+02;
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P., Longmire J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BCE45D1F31C7CE7B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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RESULT 14
Q7PB25
ID Q7PB2
AC Q7PB2
AC Q7PB2
DT 01-MA
DT 01-MA
DT 01-MA
DT 01-MA
CON Mames
OS Ricke
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                                                               Q7PB25 PRELIMINARY; PRT; 589 AA. Q7PB25, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Complete genome sequence of Bacillus thuringiensis 97-27."; Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.

-- SIMILARITY: Belongs to the ABC transporter family.

EMBL; AE017355; AAT59551.1; --
GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0005524; F:ATPase activity, coupled to transmembrane moderates.
           Multidrug resistance ABC transporter ATP-bindir
Name=rsib orf.420;
Rickettsia sibirica.
Bacteria; Proteobacteria; Alphaproteobacteria;
                                                                                                                                                                                                                                                                                                PROSITE; PS50929; ABC_TM1F; 1.
PROSITE; PS00211; ABC_TRANSPORTER 1;
PROSITE; PS50893; ABC_TRANSPORTER_2;
ATP-binding; Complete proteome.
SEQUENCE 586 AA; 64916 MW; 662AA;
                                                                                                                                                                                                                                                                                                                                                                ProDom; PD000000; ADA;
SMART; SM00382; AAA;
                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF000664; ABC membrane; 1.
Pfam; PF00005; ABC tran; 1.
ProDom; PD000006; ABC transporter;
SMART; SM00382; AAA; I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brettin T.S., Bruce D., Challacom
Hitchcock P., Jackson P., Keim P.
Richardson P., Rubin E., Tice H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OrderedLocusNames=BT9727_1789;
Bacillus thuringiensis (subsp. konkukian).
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Multidrug resistance ABC transporter, ATP-binding and permease.
Rickettsiaceae; Rickettsieae;
                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR003593; AAĀ ATPase.
InterPro; IPR011527; ABC membrane 1.
InterPro; IPR01140; ABC TM transpt.
InterPro; IPR003439; ABC transporter.
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                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                              Conservative
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Keim P., Longmire J.,
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Pred. No. 1e+02;
; Mismatches
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Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                   662AA920E116F5C4 CRC64;
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                                                   otation update)
ATP-binding protein
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              Rickettsiales
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Lucas S., O
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RESULT 15
006967
00696
AC 00696
AC 00696
DT 01-JII
DT 01-JII
DT 01-JII
DT 025-Q
GN Name
OS Bacti
OX NCBI
RN [1]
RN [2]
RP SEQ
RA Den
RL SUD
RP SEQ
RA (1)
RY SEQ
RA (2)
RP SEQ
RA (2)
RP SEQ
RA (3)
RA (2)
RA (3)
RA (4)
RA (5)
RA (6)
RA (7)
RA
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RGO; GO:0016021; C:integral to membrane; IEA.

RGO; GO:0005524; F:ATP binding; IEA.

RGO; GO:00042626; F:ATPase activity, coupled to transmembrane m. . . .

RGO; GO:0006810; P:transport; IEA.

RGO; GO:0006810; P:transport; IEA.

RINterPro; IPR0011527; ABC membrane 1.

InterPro; IPR001140; ABC_TM_transport

InterPro; IPR00140; ABC_TRANSPORTER.

RP100000; PR00004; ABC_TRANSPORTER 1.

RP100000; PS0020; ABC_TRANSPORTER 1; 1.

RP1001TE; PS00211; ABC_TRANSPORTER 1; 1.

RP1001TE; PS00893; ABC_TRANSPORTER 2; 1.

RP1001TE; PS00893; ABC_TRANSPORTER 2; 1.
             MEDLINE-98044033; PubMed=9384377; DOI=10.1038/36786;

Kunst F., Ogasawara N., Moszer I., Albartini A.M., Alloni G.,

Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,

A borriss R., Boursier L., Brans A., Braum M., Brighell S.C., Bron S.,

Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,

Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,

Denizot F., Devine K.M., Dusterhoft A., Ehrlich S.D., Emmerson P.T.,

A benizot F., Devine K.M., Dusterhoft A., Ehrlich S.D., Emmerson P.T.,

A Fritz C., Fujita M., Fujita Y., Funa S., Galizzi A., Galleron N.,

A filseppi G., Guy B.J., Haga K., Haiech J., Harwood C.R., Henaut A.,

A Jones L.-M., Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M.,

Klein C., Kobayashi Y., Koetter P., Koningstein G., Krogh S.,

Kumano M., Kurita K., Lapidus A., Lardinois S., Lauber J.,

Kumano M., Kurita K., Lapidus A., Liu H., Masuda S., Mauel C.,

Medigue C., Medina N., Mellado R.P., Mizuno M., Moestl D., Nakai S.,

Noback M., Noone D., O'Reilly M., Ogawa K., Ogiwara A., Oudega B.,
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Best Local Similarity
Matches 7; Conser
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Submitted (FEB-2003) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the ABC transporter family.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
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01-JUL-1997 (TrEMBLrel. 04, Created)
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preliminary data.
EMBL; AABWO1000001; EAA25659.1; -.
HSSP; Q9CHL8; 1MV5.
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[1]
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Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
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Best Local S
Matches 7
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RGO; GO:0016021; C:integral to membrane; IEA.

RGO; GO:0005524; F:ATP binding; IEA.

RGO; GO:00042626; F:ATPase activity, coupled to transmembrane m.

RGO; GO:0000166; F:nuclootide binding; IEA.

RGO; GO:0000166; F:nuclootide binding; IEA.

RGO; GO:0000166; F:nuclootide binding; IEA.

RGO; GO:000110; P:transport; IEA.

RINTEPPO; IPR001593; AAA ATPase.

RINTEPPO; IPR001527; ABC_membrane

RINTEPPO; IPR001140; ABC_TM transpt.

R InterPro; IPR00140; ABC_TTANSPORTER.

R Pfam; PF00064; ABC_TEAN; I.

R Pfam; PF00065; ABC_TEAN; I.

R Pfam; PF000006; ABC_TEAN; I.

R PFAM; PF000006; ABC_TEAN; I.
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Prescott A.M., Presscan E., Pujic P., Purnelle B., Rapoport G.,
Rey M., Reynolds S., Rieger M., Rivolta C., Rocha E., Roche B.,
Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter R.,
Scoffone F., Sekiguchi J., Sekowska A., Seror S.J., Serror P.,
Scoffone F., Soldo B., Sorokin A., Tacconi B., Takagi T., Takahashi H.,
Takemaru K., Takeuchi M., Tamakoshi A., Tanaka T., Terpstra P.,
Tognoni A., Tosato V., Uchiyama S., Vandenbol M., Vannier F.,
Vassarotti A., Viari A., Wambutt R., Wedler E., Weller H.,
Vassarotti A., Winters P., Wingat A., Yamamoto H., Yamane K.,
Yoshikawa H., Danchin A.,
Tohla C., Tanaka T., Zumstein E.,
Yoshikawa H., Danchin A.,
                                                                                                                                                                                                                                    PROSITE; PS50
PROSITE; PS00
PROSITE; PS50
ATP-binding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.", \ensuremath{^{"}}
                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSSP; Q9CHL8; 1MV5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nature 390:249-256(1997)
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                                                                                                                                             Similarity 7; Conserv
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PS00211; ABC_TRANSPORTER_1; 1.
PS50893; ABC_TRANSPORTER_2; 1.
                                                                                                         ATSSLDS
                                                                           ATSSLDS
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589 AA; 64519 MW; 8A15163B5698DA08
                                                                                                                                               Conservative
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GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.

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A_Geneseq_16Dec04:*
1: geneseqp1980s:*
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4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
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Gapop 10.0 ,
                                                                                                                                                                                                                                                                                                                   2105692 seqs, 386760381 residues
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49
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193.690 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	BB	ID	Description
1	49	100.0	107	ωį	AAY90824	Aay90824 520C9 hyb
2	49	•	243	N	AAW02280	0 52009
ω	49	8	243	N	AAW53170	0 52009
4	49	100.0	243	N	AAW80424	4 52009
G	49	•	243	7	ABW00716	
6	49	•	246	Ŋ	AAR39569	
7	49	100.0	267	4.	AAU04944	Aau04944 Humanised
8	49	100.0	409	4	AAU04945	Aau04945 Humanised
9	44	٠	534	N	AAR39571	Aar39571 Sequence
10	43	87.8	108	8	ADI26702	Adi26702 Mouse ant
11	41		107	8	ADE25834	
12	40		107	œ	ADE25833	Ade25833 Anti-alph
13	36	٠	9	4	AAB60407	
14	36	٠	9	4	AAB61592	Aab61592 Humanised
15	36	•	9	σ	AAU76520	Aau76520 Anti-Inte
16	36		9	v	AAE15818	
17	36		9	8	ADE71461	Ade71461 Variable
18	36	73.5	9	8	ADJ88015	-
19	36	٠	9	æ	ADN12061	Humanie
20	36	٠	9	8	ADP43336	Adp43336 Humanised
21	36	73.5	9	œ	ADS18660	Ads18660 Peptide s
22	36		89	N	AAR80080	Aar80080 Mouse der
23	36		92	N	AAW95477	Aaw95477 Mouse der
24	36	٠	107	4	AAB62083	
25	36	73.5	107	4	AAB62085	Aab62085 Humanised

WPI; 200 N-PSDB;

2000-338508/29. DB; AAA38908.

Ring DB;

(CHIR) CHIRON

CORP.

94US-00288981

The present invention describes a monoclonal antibody (MAb) (I) that binds to a human breast cancer antigen that is also bound by MAb 454C11 and 520C9 (produced hybridoma ATCC HB8484 and HB8696, respectively). All described is a hybridoma that produces (I). (I) is useful in specific binding assays, affinity purification, drug or toxin targeting, imaging, and genetic or immunological therapeutics for various cancers. The

imaging,

Also

Disclosure; Fig 13; 57pp; English.

treating cancer.

Monoclonal antibody capable of binding useful for affinity purification, drug

to human or toxin

breast cancer antigen targeting, imaging, and

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	
36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	36	
73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	73.5	
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ADR73589	ADP43324	ADP43326	ADN12052	ADN12050	ADJ88006	ADJ88004	ADE71450	ADE71452	ABR83157	ABR83156	ABR83149	ABG74710	ABG74702	ABG74718	AAE15812	AAB61581	AAB61583	AAB60396	
Adr73589	Adp43324	Adp43326	Adn12052	Adn12050	Adj88006	Adj88004	Ade71450	Ade71452	Abr83157	Abr83156	Abr83149	Abg74710	Abg74702	Abg74718	Aae15812	Aab61581	Aab61583	Aab60396	
Anti-AR a	Murine an	Humanised	Variable	Variable	Humanised	Murine mo	Murine mo	Humanise	Hu007 ant	Mu007 ant	Hu007 ant	Murine hu	Murine Mu	Murine hu	Human mAb	Murine 2C	Humanised	Mouse ant	

ALIGNMENTS

11-JAN-1985; 21-MAR-1986; 08-MAY-1988; 11-AUG-1994; Antigen binding site; immunoglobulin; cancer antigen; immunological; antibody; tumour; human; mucin; cancer; cytostatic; hybridoma; specific binding assay; affinity purification; drug targeting; toxin targeting; imaging; genetic; therapeutic. 08-FEB-1984; 07-JUN-1995; 25-APR-2000. US6054561-A. Homo sapiens. 520C9 hybridoma VL 29-AUG-2000 AAY90824; AAY90824 standard; protein; (first entry) 84US-00577976. 85US-00690750. 86US-00842476. 88US-00190778. 95US-00483749. domain SEQ ID NO:26. 107 B

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which is used in

represents a VL domain the exemplification of

derived from a 520C9 hybridoma, the present invention

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222 LOYAIFPYT 230

1 LOYALFPYT

9

Matches Query Match Best Local

Similarity 100. 9; Conservative

100.0%; S 100.0%; F ative 0;

Score 49; DB 2 Pred. No. 0.14;); Mismatches

N 0

Length 243;

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Best Local Similarity
Thes 9; Conserve
                                                          Variable heavy (VH) and variable light (VL) genes were cloned from a $20C9 hybridoma cDNA library, using probes directed toward the antibody constructed by connecting regions. A two single chain Fv (sFv) gene was constructed by connecting the VH and VL genes with a Ser rich polypeptide linker. The resulting $20C9 two sFv gene, which encodes the present sequence, was inserted into an expression vector, transformed into E. coli, and protein expression induced by the addn. of IPTG to the culture medium. A compsn. comprising a carrier and the 2 sFv protein prod. can be used for in vivo imaging, and drug targetting experiments. The 2 sFv protein prod. is a homodimer, in which both fragments target the same antigen, therefore giving greater binding avidity and longer tissue retention times, compared to individual sFv protein prod. fragments. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 107 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
29-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     520C9 anti-c-erbB-2 two single chain Fv construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Compsns. contg. antigen-targetting antibody fragment constructs - comprising dimer of single-chain Fv fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-FEB-1992;
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CREATIVE BIOMOLECULES INC
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-00831967
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Pred. No. 0.057;
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Sequence 243

RESULT 4

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AAW53170

AAW5

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XX AC AAW6

XX S200

XX 5200

XX S200

XX S9nt

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                                                                                                                    Matches
                                                                                                                                             Best
                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                          This represents the protein sequence of a 520C9 sFv' (single chain Fv) construct. This was constructed by connecting the Vh and Vl genes with a DNA sequence encoding a Serine rich linker. 520C9 is a monoclonal antibody useful in targeting c-erbB-2 antigen. This dimeric construct can be used in the methods of invention of imaging a preselected antigen expressed in a mammal. The methods are used in magnetic resonance imaging of c-erbB-2 or related antigens in cancer diagnosis. The blosynthetic constructs have enhanced properties as in vivo targetting agents in comparison with intact monoclonal antibodies or their Fab fragments. The dimeric constructs permit the in vivo targetting of an epitope on an antigen with greater apparent avidity, including greater tumour specificity, tumour localisation and tumour retention properties than that of the Fab fragment having the same CDRS as the construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antigen tumour;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                520C9 anti-c-erbB-2 sFv' dimeric construct protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW53170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-FEB-1992;
07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US5753204-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; AAV21798.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oppermann H,
                                                                                                                                                                                                                                  Sequence 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Col 33-36; 30pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-311318/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Imaging of antigens in vivo - using
                                                                                                                                                Local
   222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          imaging; single chain Fv; sFv; linker; dimeric; cancer; c-erbB-2;
diagnosis; ss.
                                                       1 LQYAIFPYT 9
                                                                                                                 Similarity 9; Conserv
   LQYAIFPYT 230
                                                                                                              100.0%;
llarity 100.0%;
Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry
                                                                                                                                                                                                                                      A
A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92US-00831967.
93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-00461838.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Huston JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                       0
                                                                                                                                                   Score 49; DB 2
Pred. No. 0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B
                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dimers of single-chain antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Houston LL;
                                                                                                                                                                              DB 2;
                                                                                                                          0
                                                                                                                                                                              Length 243;
                                                                                                                          Indels
                                                                                                                             0,
                                                                                                                                Gaps
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AAW80424

ID AAW80424 standard; pi
XX

AC AAW80424;

XX

AC AAW80424;

XX

DT 28-JAN-1999 (first 6
XX

DE 520C9 sFv sequence.

XX

Synthetic.

XX

Synthetic.

XX

Synthetic.

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OS Synthetic.

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OS Synthetic.

XX

OS Synthetic.

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OS Synthetic.

XX

IN-NOV-1998.

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OS SYNTHETICE SING CORP.

XX

OS SYNTHETICE BIOLOGOME.

XX

OFF 05-JUN-1995; 95US-(
YEAR 06-FEB-1992; 92US-(
YEAR 07-OCT-1993; 93US-(
YEAR 07-OCT-19
ARBSULT 5
ARBOOT16
ID ABMO
XX ABMO
XX ABMO
XX DE 520C
XX Cell
KW Cell
KW GPV
XX GPV
XX Unid
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XX Unid
XX Unid
XX Us20
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents an antibody 520C9 sFV. Variable heavy and light sequences of antibody 7520C9 are connected, together with a serine linker, to produce the present single chain Fv gene. The present sequence exemplifies the invention. Dimers of the single chain Fv are used for targeted delivery of drugs or imaging agents (e.g. cytotoxins, prodrugs or 99m-technetium) to antigen-expressing cells, particularly for treatment or diagnosis of tumours (especially of ovary or breast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid encoding single-chain Fv fragment specific for antigens - and having C-terminal tail for crosslinking to form dimer with improved pharmacokinetic properties, used to deliver drugs and imaging agents, especially to tumours.
                                                                                                                                                                                            520C9 sFv protein
                                                                                                                                                                                                                                                         15-JAN-2004
                                                                                                                                                                                                                                                                                                              ABW00716;
US2002168375-A1
                                                                                                                                                                                                                                                                                                                                                                   ABW00716 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-023541/02.
N-PSDB; AAV63399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Col 35-36; 29pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antigen-expressing cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                520C9 sFv; antigen; tumour cell; antibody 520C9; targeted delivery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CREA-) CREATIVE BIOMOLECULES INC (CHIR ) CHIRON CORP.
                                                                                                                                       proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 LQYAIFPYT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOYAIFPYT 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
llarity 100.0%;
Conservative 0;
                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-00831967.
93US-00133804.
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                                                                                                                                    cytotoxic agent; drug targetting; single-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Huston JS,
                                                                                                                                                                                                                                                                                                                                                                      243 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 49; DB 2;
Pred. No. 0.14;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 243;
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ARBSULT 6
AAR3956
ID AAR39
XX AAR3
AC AAR3
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DT 25-M
DT 07-F
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KW Sing
KW bios
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Matches
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07-OCT-1993;
05-JUN-1995;
26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a binding protein formulation for targetting an epitope on an antigen expressed in mammal. The formulation comprises of dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain. The invention is useful for imaging a preselected antigen in a mammal expressing the antigen. It is also useful for targetting drugs that inhibits cell proliferation and cytotoxic agents that kill cells. The present sequence is 52009 sfv protein. This sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                  Single chain biosynthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
07-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel binding protein formulation for targeting epitope on antigen in mammal, comprising dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain.
                          Huston JS,
                                                                                                                                                  06-FEB-1992;
                                                                                                                                                                                                 05-FEB-1993;
                                                                                                                                                                                                                                                     19-AUG-1993
                                                                                                                                                                                                                                                                                                       WO9316185-A2
                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence of 520C9 sFv protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR39569;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR39569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 19-20; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-765156/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Huston JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CHIR ) CHIRON CORP
                                                                         (CREA-) CREATIVE BIOMOLECULES (CETU) CETUS ONCOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 LOYALFPYT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 100 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD61485.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; protein; 246 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ||||||||
LQYAIFPYT 230
                          Houston LL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ŗ,
                                                                                                                                                                                                                                                                                                                                                                                                  Fv polypeptide; VH-VL heterodimer; immunoglobulin; single polypeptide chain binding site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-00558741
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92US-00831967.
93US-00133804.
95US-00462641.
                                                                                                                                                  92US-00831967
                                                                                                                                                                                                   93WO-US001055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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                          Ring
                                                                                                INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 49; DB 7
Pred. No. 0.14;
                          BG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                          Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 7;
                          Ξ,
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RESULT 7
AAU04944
ID AAU0
XX AAU0
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DT 06-A
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XX Huma
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OS Mus
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Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. such as breast and ovarian tumour cells, which is an approx.
200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). A single chain Fv (sFv) is a covalently linked VH-VL heterodimer which is expressed from a gene fusion including VH- and VL- encoding genes connected by a peptide-encoding linker. Such linker sequences are set forth in AA residues 116-135 in AAR39569, which includes part of the 16 AA linker sequences in AAR39572. Using AAQ46084 for the 520C9 monoclonal antibody, a single chain polypeptide can be produced having a binding affinity for a c-erbB-2 related antigen. 'X' in AAR39569 refers to the location of a stop codon in AAQ46084. (Updated on 55-MAR-2003 to correct PN field.)
                                                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                                                                                                                                                     Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Humanised; antibody; p185; 520C9H; interleukin-2; IL-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU04944;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ds sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Humanised anti-p185 single chain antibody, 520C9H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAU04944 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New single chain Fv imaging or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 60-61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAQ46084.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1993-272889/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOYAIFPYT 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOYAIFPYT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adenocarcinoma.
                                                                                                        /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                             /label= CDR
/note= "Complementarity determining region"
                                                  /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                      'label= Synthetic_peptide_linker
'note= "Links the heavy chain to
                                                                                                                                                                                                                            /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                    note= "Complementarity determining region'
                      label=
                                                                                                                                                                                                                                                                                                            .abel= CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polypeptide binding to C-erbB-2 tumour antigen - breast or ovarian cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                              .208
                                                                                                                                                                                                                                                                    .126
                                        .249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              87pp; English.
                                                                                                                                                                                                              .152
                                                                                                                                                        .186
                        CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 49;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 246;
                                                                                                                                                                      the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                    light chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunoconjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>,</u>
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RESULT 8
AAU04945
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Matches
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                                                                                                                                                             Mus sp.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence represents a humanised anti-p185 single chain antibody which is linked to a human interleukin-2 (II-2) molecule to make a fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cancers particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, renal system, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
 Region
                                                                                                                Key
                                Protein
                                                                Protein
                                                                                                Peptide
                                                                                                                                              Chimeric.
                                                                                                                                                                                                                              cancer;
                                                                                                                                                                                                                                               Humanised;
                                                                                                                                                                                                                                                                             Humanised
                                                                                                                                                                                                                                                                                                            06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                             AAU04945;
                                                                                                                                                                                                                                                                                                                                                                                          AAU04945 standard; protein; 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 7; Fig 9; 74pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel immunoconjugate comprises a humanized
                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Austin R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR )
(HAMI-)
(HAMI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JAN-2000; 2000US-0177258P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JAN-2001; 2001WO-US001919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-JUL-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            241
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          μ
                                                                                                                                                                                                                            tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
9; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HAMILTON CIVIC HOSPITALS HAMILTON REGIONAL CANCER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS09507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LQYAIFPYT 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOYAIFPYT 9
                                                                                                                                                                                                                                                                            anti-p185 antibody/IL-2 fusion protein
                                                                                                                                                                                                                           antibody; p185; 520C9H; interleukin-2;
mour; adenocarcinoma; fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                          (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA;
/label= Signal_peptide
23. .409
/label= Mature_fusion_protein
23. .259
/label= Humanised_antibody_520C
53. .57
                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        260. .267
/label= Glu_Glu_epitope
                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for inhibiting tumor cell growth in vivo anti-pl85 antibody linked to an Interleukin-2
             Humanised_antibody_520C9H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 49;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 267;
                                                                                                                                                                                                                                            IL-2;
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                                                                                                                                                                                                                                            immunoconjugate;
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                                                                                      Query Match
Best Local S
Matches
                                                                                                         The sequence represents a humanised anti-p185 (520C9) single chain antibody/human interleukin-2 (IL-2) fusion protein. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cances particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, rensestem, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                     Novel immunoconjugate useful for inhibiting tumor cell comprises a humanized anti-pl85 antibody linked to an I polypeptide.
                                                                                                                                                                                                                                                                                                                          (CHIR )
(HAMI-)
(HAMI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Region
                                                                                      Sequence 409
                                                                                                                                                                                                Claim 13; Fig 11; 74pp; English
                                                                                                                                                                                                                                                                                                    Austin R, Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                                              19-JAN-2001; 2001WO-US001919
                                                                                                                                                                                                                                                                                                                                                                                                                   26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200153354-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Region
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                                                                                                                                                                                                                                                                                                                                                                       20-JAN-2000; 2000US-0177258P
                                                                                                                                                                                                                                                                    2001-451904/48
DB; AAS09508.
241
                                                                                                                                                                                                                                                                                                                           CHIRON CORP.

HAMILTON CIVIC HOSPITALS
HAMILTON REGIONAL CANCER
                                           Similarity
9; Conserv
                      LQYAIFPYT 9
LOYAIFPYT 249
                                           Conservative
                                                                                      AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              'note= "Encode by ACA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= IL_2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= Synthetic_linker_peptide
note= "Links the antibody to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          label= CDR note= "Complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           label= CDR
note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           label= CDR note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            label- Synthetic_peptide_linker
note- "Links the heavy chain to the light chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              label- CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "Encoded by GGG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 label=
                                      100.0%; 5C
100.0%; Pr
                                                                                                                                                                                                                                                                                                    Ring
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 152
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                                                   Score 49; DB 4
Pred. No. 0.23;
                                           Mismatches
                                                                                                                                                                                                                                                                                                                             CENT
                                                                                                                                                                                                                                                                                                                                      RES DEV INC.
                                                               DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the IL-2 molecule"
                                                               Length 409;
                                           Indels
                                                                                                                                                                                                                                   Interleukin-2
                                          <u>,</u>
                                          Gaps
                                                                                                                                                      cancers
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RESULT 9

02-OCT-2003.

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                                                                                                                                                                                       RESULT 10
                                                                                                                                                                                                                                                                 Query Match
Best Local &
                                                                                                                                                                                                                                                          Matches
                                                             axon; glial cell proliferation;
Theiler's murine encephalomyelitis virus infection; CNS injury;
                                                                                 Mouse; antibody; IgM; remyelination; neuronal growth; autoantibody; demyelination disease; multiple sclerosis; central nervous system; CNS;
                                                                                                                                                                                                                                                                                                                        c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells, which is an approx. cells, which is an approx. 200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                         New single chain Fv polypeptide binding to C-erbB-2 tumour antigen - imaging or treating breast or ovarian cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumour antigen; c-erbB-2; G-FIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
07-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR39571 standard; protein;
                  US2003185827-A1
                                    gs snw
                                                                                                            Mouse anti IgM antibody HNK-1 kappa light chain protein.
                                                                                                                                                  ADI26702;
                                                                                                                                                                    ADI26702 standard; protein; 108
                                                                                                                                                                                                                                                                                              Seguence 534 AA;
                                                                                                                                                                                                                                                                                                                                                                       Example; Page 65-68; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1993-272889/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Huston JS, Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CREA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-FEB-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9316185-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence of G-FIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR39571;
                                                                                                                                15-APR-2004
                                                                                                                                                                                                                    513
                                                       cord
                                                                                                                                                                                                                                     μ
                                                                                                                                                                                                                                                        8; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ46086.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CREATIVE BIOMOLECULES CETUS ONCOLOGY CORP.
                                                                                                                                                                                                                    LQYPIFPYT 521
                                                                                                                                                                                                                                    LOYALFPYT
                                                     injury.
                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(first ent
                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                   89.8%;
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                                                                                                                                                                                                                                                        Score 44; DB
Pred. No. 3.2;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB,
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3.2;
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                                                                                                                                                                                                                                                                         Length 534
                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                     of about
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89 LQYASFPYT 97

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cc mammal (comprising administering a monoclonal antibody, or mixtures, contrained by their ability to bind structures and cells within the colls in CNS, including oligodendrocytes), stimulating the proliferation of gilal cells in CNS axons in a mammal (comprising administering a monoclonal colls within the colls in CNS axons in a mammal (comprising administering a monoclonal colls within the colls in CNS axons in a mammal (comprising administering a monoclonal colls within the CNS), treating or preventing a monoclonal colls within the CNS), treating or preventing a monoclonal colls structures and cells within the CNS) in the comprising administering a considering a monoclonal colls within the CNS in a mammal (comprising administering a composition of colls in the CNS), treating or preventing a composition of colls in the CNS, in a mammal (comprising administering a composition of colls in the CNS), treating or preventing a composition of colls in the colls within the CNS, and to stimulate remyelination of colls and colls within the CNS, and to stimulate remyelination of colls axons of the CNS, stimulating, in vitro, the proliferation of gilal colls within the CNS, and to stimulate remyelination of CNS axons in composition of colls axons of the CNS, and to stimulate remyelination of colls axons of colls and the colls within the colls and to stimulate remyelination of colls axons of colls and other agents for the ability to modulate the production or mimic the colls and to stimulate the proliferation of the CNS comprising using colls colls in the colls and to stimulate the proliferation of the CNS comprising using colls colls in the colls colls in CNS axons, collected with strain DA of Theiler's collected with strain DA of Theiler's collected colls colls in cNS axons, composition of the CNS, the colls of the colls of the colls of the colls c
                       Query Match
Best Local
      Matches
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30-MAY-2000;
05-DEC-2000;
                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunocompetent host with an antibody peptide, and harvesting the antibody, where the peptide comprises a human anti-1gM antibody fragm given in the specification, or active fragments. Also included are stimulating remyelination of central nervous system (CNS) axons in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human immunoglobulin M antibody for treating or preventing a demyelinating disease of the central nervous system in a human or domestic animal, such as multiple sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-AUG-1996;
07-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an antibody (I) produced by injecting an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 19; Fig 69; 159pp; English.
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  Similarity
8; Conserv
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    Conservative
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; 96US-00692084.
; 97US-00779784.
; 99US-00322862.
; 2000US-00580787.
; 2000US-00730473.
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                       88.9%;
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CC comprising heavy chain complementarity determining regions (CDR) 1, 2, CC and 3, or a heavy and light chain variable domain sequence; (2) a CC monoclonal antibody that specifically binds to alpha-v-beta-6 but does CC not inhibit binding of alpha-v-beta-6 to LAP; (3) a composition for comprising the antibody and a carrier; (4) a method for treating a CC comprising the antibody and a carrier; (4) a method for treating a Subject having or a trisk of having a disease mediated by alpha-v-beta-6 in a mammal CC preventing or postponing the onset of the disease; (5) a method of CC alleviating or postponing the onset of the disease; (5) a method of CC detecting alpha-v-beta-6 in a tissue sample by contacting the tissue CC sample with the antibody; and (6) a cell hybridoma 6.LAB, 6.3G9, 6.8G6, 6.2B1, 7.1G10, 7.7G5, or 7.1Cr, which respectively comprises American CC Type Culture Collection (ATC) Accession number PTA-3640, PTA-3649, PTA-3899, PTA-3899, or PTA-3900. (I) has cytostatic, CC and can be used in vaccines. The antibodies, compositions and methods of the present invention can be used for preventing or treating a disease mediated by alpha-v-beta-6, e.g. fibrosis such as scleroderma, scarring, CC liver fibrosis, kidney fibrosis or lung fibrosis; psoriasis; cancer, or Layout a syndrome. The present sequence is used in the exemplification of the present invention. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a monoclonal antibody (I) that specifically binds to alpha-v-beta-6, and inhibits the binding of -beta-6 to latency associated peptide (LAP) with an ICSO value low that of 1005. Also described: (1) an anti-alpha-v-beta-6 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New monoclonal antibody that specifically binds to alpha-v-beta-6, and inhibits the binding of alpha v beta 6 to latency associated peptide (LAP), useful for treating fibrosis, psoriasis, cancer, or Alport's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 27; SEQ ID NO 41; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 syndrome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic; dermatological; vulnerary; hepatotropic; immunosuppressive; vaccine; fibrosis; scleroderma; scarring; liver fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAR-2002;
13-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-MAR-2003; 2003WO-US008048
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             kidney fibrosis; lung fibrosis; psoriasis; cancer; Alport's syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-alpha-v-beta-6 monoclonal Ab light chain variable domain SEQ ID:41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     monoclonal antibody; alpha-v-beta-6; latency associated peptide; LAP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-2004
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2002US-0426286P.
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DB 8; Length 107;

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         monocional antibody that specifically binds to alpha-v-beta-6 but does not inhibit binding of alpha-v-beta-6 to LAP; (3) a composition for preventing or treating a disease mediated by alpha-v-beta-6 in a mammal comprising the antibody and a carrier; (4) a method for treating a by administering to the subject the composition described above, and so alleviating or postponing the onset of the disease (5) a method of detecting alpha-v-beta-6 in a tissue sample by contacting the tissue sample by contacting the tissue sample by contacting the tissue sample by the composition of the composition and method of the present invention can be used for preventing or treating a disease mediated by alpha-v-beta-6, e.g. fibrosis such as scleroderma, scarring, liver fibrosis, portals; portals; portals; portals; procession contents or lung fibrosis, portals; cancer,
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Best Local
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                                                                                                                                                                                                                                                                                                                    specifically binds to alpha-v-beta-6, and inhibits the binding of alph-beta-6 to latency associated peptide (LAP) with an IC50 value lower that of 1005. Also described: (1) an anti-alpha-v-beta-6 antibody comprising heavy chain complementarity determining regions (CDR) 1, 2, and 3, or a heavy and light chain variable domain sequence; (2) a
                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a monoclonal antibody (I) that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      monoclonal antibody; alpha-v-beta-6; latency associated peptide; LAP; cytostatic; dermatological; vulnerary; hepatotropic; immunosuppressive; vaccine; fibrosis; soleroderma; scarring; liver fibrosis; kidney fibrosis; lung fibrosis; psoriasis; cancer; Alport'; syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 26; SEQ ID NO 40; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New monoclonal antibody that specifically binds to alpha-v-beta-6, and inhibits the binding of alpha v beta 6 to latency associated peptide
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13-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (LAP), useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    68
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cancer, oral,
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1; Mismatches
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RESULT 13
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CC comprising administering an antibody which binds ErbB2 (HER; AAB60408).

CC In particular, the anti-ErbB2 antibody is the murine monoclonal antibody CC 2C4 (AAB60396, AAB60397) or a humanised version of 2C4 (AAB60398). The invention also encompasses an isolated nucleic acid comprising such nucleic acids; the recombinant production of a humanised CC comprising such nucleic acids; the recombinant production of a humanised CC erbB2-binding antibody; and an immunoconjugate comprising a humanised CC erbB2-binding antibody and a cytotoxic drug. The ErbB2-binding antibodies act by antagonising ErbB receptors, and as inhibitors of transforming CC growth factor alpha (TGF-alpha)-activated mitogen activated protein CC kinase (MAPK). The method of the invention is used for treating cancer, cespecially colon cancer, rectal cancer, colorectal cancer, lung cancer (especially cono-small cell lung cancer), or breast cancer (especially non-small cell lung cancer), or breast cancer (especially corest cancer). The antibodies may also have non-therapeutic cuses e.g., as affinity purification agents. Using an antibody which binds cto ErbB2 to treat cancer is preferable to the use of EGFR targetted concer, as EGFR is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is also highly expressed in other tissues such as the colored concer is a colored concer is colored concer is colored to colored concer is also highly expressed in other tissues such as the colored concer is colored to colored concer is colored to colored concer is colored to co
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-ErbB2 monoclonal antibody; HER2; humanised; VL; light chain variable region; cancer; cytostatic; EGFR-expressing cancer; epidermal growth factor receptor; colon cancer; rectal cancer; tumour; colorectal cancer; non-small cell lung cancer; metastatic breast cancer; affinity purification; complementarity determining region; CDR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 107
                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method for treating cancer in a human patient, wherein the cancer expresses epidermal growth factor receptor (EGFR),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating cancer in a human, where factor receptor (EGFR), comprises
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 66; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Presta LG,
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77.8%;
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Pred. No.
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                                                                                                                       The ErbB family of receptor tyrosine kinases are important mediators of cell growth, differentiation and survival. The receptor family includes four distinct members including Epidermal Growth Factor Receptor (EGFR or ErbB1), HER2 (ErbB2 or p185 neu), HER3 (ErbB3) and Her4 (ErbB4 or tyro2). The present invention relates to a method for treating prostate cancer. The method comprises administering an antibody which binds ErbB2 and blocks ligand activation of an ErbB receptor. Preferably, the antibody blocks binding of monoclonal antibody 2C4 to ErbB2 and/or blocks TGF-alpha activation of mitogen-activated protein kinase (MAPK). The present sequence is a complementarity determining region (CDR) from the variable light chain of humanised Fab version 574 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                    Treating prostate cancer in a human which binds ErbB2 and blocks ligand
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                                                                                                                                                                                                                                                                                                       Disclosure; Page 24; 93pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                     (SETH ) GENENTECH INC.
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Matches 6; Conserv
                                                                                                                                                                                        The invention relates to novel isolated mammalian anti-interleukin-12 (II -12) antibodies. The antibodies comprise at least one complementarity determining region (CDR) of a heavy or light chain, a heavy chain or light chain variable region, or a heavy chain or light chain constant region. The anti-II-12 antibodies are useful in methods for diagnosing or treating II-12 related conditions, e.g. psoriasis or multiple sclerosis. The antibodies are also useful for treating immune, cardiovascular, infectious, malignant or neurological disorders or diseases. The present sequence represents the amino acid sequence of human anti-Interleukin- 12 (II-12) antibody CDR3 light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-AUG-2000; 2000US-0223358P.
29-SEP-2000; 2000US-0236827P.
01-AUG-2001; 2001US-00920262.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; antibody; anti-interleukin-12; CDR; light chain; circulatory; complementarity determining region; neuroprotective; antitipsoriatic; immunostimulant; cytostatic; anti-microbial; psoriasis; infection; multiple sclerosis; immune disorder; cardiovascular; malignant disease;
                                                                                                                                                                                                                                                                                                                                                                                         New mammalian anti-IL-12 antibodies, useful for diagnosing or treating -12 related conditions, e.g. psoriasis or multiple sclerosis, as well enther for treating immune, infectious, malignant or neurological
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Copyright (c) 1993 - 2005 Compugen Ltd.
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/ Cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
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/ Cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
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/ Cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
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US-10-912-994-6
US-10-437-963-137213
US-10-268-501-1
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US-09-766-543-10
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        Sequence 6, Appli
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Sequence 145, Appl
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	Sequence 246038,	O	æ	Φ		Sequence 9, Appli	6, Appl	о 2	2, Appli	e 7:	5	œ	۲,	_	270138	7392	e 10, Ap	Sequence 5, Appli	10,	e 1	ູ້	4	о 8	Sequence 8, Appli	e U	e u	æ	e 12, App	e 10, App	equence 3, Appl		

ALIGNMENTS

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US-09-887-853-6
US-09-887-853-6
J Sequence 6, Application US/09987853
Patent No. US20020168375A1
J PATENTI INFORMATION:
GENERAL INFORMATION:
HOUSTON, L. L.
Ring, David B.
RITLE OF INVENTION: Biosynchetic Binding Proteins For Invention: Exchange Place, 53 State Street STREET: Exchange Place, 53 State Street CITY: Boston STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
APPLICATION NUMBER: US/09/887,853
FILING DATE: 21-Unn-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATE: CUNknown>
PRIOR APPLICATION DATA:
APPLICATION DATE: CUNknown>
ATTORNEY/AGENT INFORMATION:
NAME: Kalley, Robin D.
REGISTRATION NUMBER: 34,637
REFERRENCE/DOCKET NUMBER: 34,637
REFERRENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
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US-09-887-853-6
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TITLE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING
FILE REFERENCE: CIBT-P01-130
CURRENT APPLICATION NUMBER: US/10/683,547
CURRENT FILING DATE: 2003-10-10
PRIOR PPLICATION NUMBER: US/09/558,741
PRIOR FILING DATE: 2000-04-26
PRIOR FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 07/831,967
PRIOR FILING DATE: 1992-02-06
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
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Patent No. US20020041865A1
GENERAL INFORMATION:
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SEQ ID
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                                                   TITLE OF INVENTION: METHODS FOR TREATING TUMORS FILE REFERENCE: PP01679.002 CURRENT APPLICATION NUMBER: US/09/766,543 CURRENT FILING DATE: 2000-01-19 PRIOR APPLICATION NUMBER: 60/177,258 PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                    APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Ring, David B.
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TELEPAX: 617-248-7100
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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ORGANISM: Artificial Sequence
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mes 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                              222
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                                                                                                                                                                                                                                                                                                                                                                                                LOYAIFPYT 9
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TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 49; DB 1 100.0%; Pred. No. 0.27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Kwok, Cheuk S.
APPLICANT: Ring, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PD16679.002
CURRENT APPLICATION NUMBER: US/09/766,543
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
NUMBER: OF SEQ ID NOS: 14
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US-10-010-729-45
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LENGTH: 276
TYPE: PRT
                                                                                                                                           Sequence 45, Application US/10010729

Publication No. US20030185827A1

GENERAL INFORMATION:

APPLICANT: Rodriguez, Moses
APPLICANT: Miller, David J.

APPLICANT: Pease, Larry R.

TITLE OF INVENTION: Therapeutic Uses Thereof Particularly:

TITLE OF INVENTION: System

FILE REFERENCE: 1199-1-005CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
           CURRENT APPLICATION NUMBER: US/10/010,729
CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 09/730,473
PRIOR FILING DATE: 2000-12-05
PRIOR PILING DATE: 2000-12-05
PRIOR APPLICATION NUMBER: 09/580,787
PRIOR APPLICATION NUMBER: 09/322,862
PRIOR APPLICATION NUMBER: 09/322,862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: 520C9 OTHER INFORMATION: humanized single-chain antibody used in the OTHER INFORMATION: IL-2-antibody fusions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: Description of Artificial Sequence: H520C9sFv plus OTHER INFORMATION: linker
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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les 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                   241 LQYAIFPYT 249
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1999-05-28
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Pred. No. 0.31;
Mismatches
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Pred. No. 0.
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FILING DATE:

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US-09-920-262A-6
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US-10-437-963-119518
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; TYPE: PRT
; ORGANISM: Mus musculus
US-10-010-729-45
GENERAL INFORMATION:

APPLICANT: Shealy, David

APPLICANT: Shealy, David

APPLICANT: Scallon, Bernie

APPLICANT: Giles-Komar, Jill

APPLICANT: Giles-Komar, Jill

APPLICANT: Parit, David

TITLE OF INVENTION: ANTI-11-12 ANTIBODIES, COMPOSITIONS, METHODS AND USES

FILE REFERENCE: CENO248

CURRENT APPLICATION NUMBER: US/09/920,262A

CURRENT PILING DATE: 2002-05-06

PRIOR APPLICATION NUMBER: 60/223,358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(5221)B CURRENT APPLICATION NUMBER: US/10/437,963 CURRENT FILING DATE: 2003-05-14 NUMBER OF SEQ ID NO 119518 SEQ ID NO 119518 LENGTH: 217
                                                                                                                                                                                                                                   Sequence 6, Application US/09920262A Publication No. US20030124123A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 119518, Application US Publication No. US20040123343A1 GENERAL INFORMATION:
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version SEQ ID NO 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1997-01-07
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR FILING DATE: 1996-08-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 08/236,520 PRIOR FILING DATE: 1994-04-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Oryza sativa FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
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Similarity 85.7%;
6; Conservative
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Similarity 88.9%;
8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/10437963
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; ORGANISM: Mus musculus US-10-608-626-12
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US-10-608-626-12
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US-10-268-501-12
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                                                                                                                                                                                                                                 Sequence 12, Application US/10608626

Publication No. US20040013667A1

GENERAL INFORMATION:

APPLICANT: Kelsey, Stephen M.

APPLICANT: Sliwkowski, Mark X.

TITLE OF INVENTION: Treatment with Anti-BrbB2 Antibodies

FILE REFERENCE: P1467R2P2

CURRENT APPLICATION NUMBER: US/10/608,626

CURRENT FILING DATE: 2003-06-27

CURRENT FILING DATE: 2003-06-27
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                                                                            SEQ ID NO 12
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TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies
FILE REFERENCE: P1467R2P1
CURRENT APPLICATION NUMBER: US/10/268,501
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR FILING DATE: 2000-06-23
PRIOR FILING DATE: 1999-06-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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PRIOR APPLICATION NUMBER: 60/236,827
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver 3.1
                                                                                                               PRIOR APPLICATION NUMBER: US 10/268,501
PRIOR FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 1999-06-25
                                                                                               NUMBER OF SEQ ID NOS: 13
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TYPE: PRT
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                                       LENGTH: 9
TYPE: PRT
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ORGANISM: Homo sapiens
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Local Similarity 75.0%;
les 6; Conservation
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Local Similarity 75.0%;
les 6; Conservative
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No. US20030086924A1
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Pred. No. 1.3e+06;
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Query Match
Best Local Similarity
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_38717C.1.pep US-10-437-963-137213
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US-10-437-963-137213
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TITLE OF INVENTION: ANTI-11-12 ANTIBODIES, CC
TITLE REFERENCE: CEN0248DIY1
CURRENT APPLICATION NUMBER: US/10/912,994
CURRENT FILING DATE: 2004-08-06
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR FILING DATE: 2001-08-01
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                                                                                                    APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(52921)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 137213
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 137213, Application US/10437963
Publication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver 3.1 SEQ ID NO 6
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APPLICANT: Knight, David
APPLICANT: Peritt, David
APPLICANT: Scallon, Bernie
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TYPE: PRT
ORGANISM: Homo sapiens
                                     TYPE: PRT
ORGANISM: Oryza sativa
FEATURE:
                                                                                                ENGTH: 88
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Pred. No. 1.3e+06;
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Pred. No. 1.3e+06;
1; Mismatches 1; Indels
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                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 199-06-25
NUMBER OF SEQ ID NOS: 13
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/10268501 Publication No. US20030086924A1 GENERAL INFORMATION:
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APPLICANT: Sliwkowski, Mark X.
TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies
FILE REFERENCE: P1467R2P1
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Best Local Similarity 75.0
Matches 6; Conservative
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Publication No. US20030086924A1
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Best Local S
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TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies
FILE REFERENCE: P1467R2P1
CURRENT APPLICATION NUMBER: US/10/268,501
CURRENT FILING DATE: 2002-10-10
CURRENT FILING DATE: 2002-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/268,501
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 13
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TYPE: PRT
ORGANISM: Mus Musculus
                                                                                                                                                                FEATURE:
                                                                                                                                                                             ORGANISM: Artificial sequence
                                                                                                                                                                                               TYPE: PRT
                                                                                                                                                                                                              LENGTH: 107
90
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6; Conserva
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QYYIYPYT 97
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                                                              Conservative
                                                                               73.5%;
75.0%;
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                                                               Score 36; DB Pred. No. 34; 1; Mismatches
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Pred. No. 34;
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RESULT 14

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RESULT 15

US-10-608-626-3

i Sequence 3, Application US/10608626

publication No. US20040013667A1

GEMERAL INFORMATION:

APPLICANT: Kelsey, Stephen M.

APPLICANT: Sliwkowski, Mark X.

ITTLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies

FILE REFRENCE: P1467R2P2

CURRENT APPLICATION NUMBER: US/10/608,626

CURRENT FILING DATE: 2003-06-27

PRIOR APPLICATION NUMBER: US 10/268,501

PRIOR FILING DATE: 2002-10-10

PRIOR FILING DATE: 2002-06-23

PRIOR APPLICATION NUMBER: US 09/602,812

PRIOR PILING DATE: 1999-06-25

NUMBER OF SEQ ID NOS: 13

SEQ ID NO 3

LENGTH: 107

TYPE: PRT

ORGANISM: Artificial Sequence

PEATURE:
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APPLICANT: Kelbey, Stephen M.

APPLICANT: Kelbey, Stephen M.

TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies

FILE REFERENCE: P1467R2P2

CURRENT APPLICATION NUMBER: US/10/608,626

CURRENT APPLICATION NUMBER: US 10/268,501

PRIOR APPLICATION NUMBER: US 9/602,812

PRIOR FILING DATE: 1999-06-25

NUMBER OF SEQ ID NOS: 13
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Search completed: April 4, 2005, 16:42:45 Job time : 13.2843 secs
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; OTHER INFORMATION: humanized VL sequence
US-10-608-626-3
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; TYPE: PRT
; ORGANISM: Mus Musculus
US-10-608-626-1
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Best Local Similarity
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RESULT 2
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Ig kappa chain V region - mouse

C;Species: Mus musculus (house mouse)

C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 21-Jan-2000
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R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson, Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.; Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L. Nature 390, 364-370, 1997

Nature 390, 364-370, 1997

A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artlach, P.; Kaine, B.P.; Sykes, S.N. Smith, H.O.; Woese, C.R.; Venter, J.C.

Smith, H.O.; Woese, C.R.; Venter, J.C.

Smithe: The complete genome sequence of the hyperthermophilic, sulfate-reducing archaeor A;Reference number: A69250; MUID:98049343; PMID:9389475

A;Accession: F69371
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F69371
bile acid-inducible operon protein F (baiF-2) homolog - Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Species: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
                                                                                                         A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-106 <LOS>
A;Cross-references: EMBL:X65008; NID:g52649; PIDN:CAA46141.1; PID:g52650
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;16-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                   C;Accession: $20652
R;Losman, M.; Fasy, T.M.; Novick, K.E.; Monestier, M. submitted to the EMBL Data Library, February 1992
A;Description: Relationships among antinuclear antibodies A;Reference number: $20639
A;Accession: $20652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA A;Residues: 1-450 <KLE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:029288; GB:AE001037; GB:AE000782; NID:g2689360; PIDN:AAB9026
Matches
                                                          Query Match
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Local Similarity 85.7%;
les 6; Conservation
                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                275
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  Similarity 77. 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                YAVFPYT 281
                          73.5%;
  <u>,</u>
                             Score 36; DB Pred. No. 5.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 39;
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  Mismatches
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                                                          DB 2;
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                                                       Length 106;
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     Indels
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  Gaps
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  0;
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Ig light chain V region (clone 202.105) - mouse (fragment)
(;Species: Mus musculus (house mouse)
(;Date: 30-Sep-193 #sequence_revision 30-Sep-1993 #text_change 21-Jan-2000
(;Accession: PH1062
R;fillman, D.M.; Jou, N.T.; Hill, R.J.; Marion, T.N.
J. Exp. Med. 176, 761-779, 1992
A;Title: Both IgM and IgG anti-DNA antibodies are the products of clonally selective
A;Reference number: PH0971; MUID:92381444; PMID:1512540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein SPAC8A4.04c - fission yeast (Schizosaccharomyces pombe) (;Species: Schizosaccharomyces pombe C;Date: 16:May-1996 #sequence_revision 13-Mar-1997 #text_change 31-Jan-2000 C;Accession: T3912c; S62520 R;Lye, G.; Churcher, C.M.; Barrell, B.G.; Rajandream, M.A.; Walsh, S.V. submitted to the EMBL Data Library, November 1995 A;Reference number: Z21830 A;Reference number: Z21830 A;Accession: T39126 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-177 - LY2> A;Residue
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C; Genetics:
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S62520
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A;Residues: 1-107 <BUL>
A;Cross-references: UNIPROT:Q57895; GB:U67496; GB:L77117; NID:g2826283; PIDN:AAB98442.1;
C;Genetics:
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A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.; Woese, A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschi: A;Reference number: A64300; MUID:96337999; pMID:8688087
A;Accession: E64356
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E64356
                                                                                                                                                                                                                                                                                                                               RESULT 5
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C;Superfamily: Uncharacterized conserved protein
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85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 36; DB
Pred. No. 8.4;
2; Mismatches
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Pred. No. 5.1;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8.4;
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                                               В
   RESULT 8
PL0262
Ig kappa chain V region (anti-DNA, 60VK) - mouse (fragment)
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C28840
Ig kappa chain V region (HP22) - mouse (Iragiment,
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 29-Aug-1987 #sequence_revision 29-Aug-1987 #text_change 21-Jan-2000
C;Accession: C28840; J25114
R;Ollier, P:, Rocca-Serra, J:, Somme, G.; Theze, J.; Fougereau, M.
EMBO J. 4, 3681-3688, 1985
A;Title: The idiotypic network and the internal image: possible regulation of a germ-line
A;Reference number: A91028; MUID:86136012; PMID:3937730
                                                                                                                                               C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;10-84/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                             Ig kappa chain V region (HP27) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 29-Aug-1987 #sequence_revision 29-Aug-1987 #text_change 21-Jan-2000
C;Accession: B28840; IZ5114
R;Ollier, P.; Rocca-Serra, J.; Somme, G.; Theze, J.; Fougereau, M.
EMBO J. 4, 3681-3688, 1985
A;Title: The idiotypic network and the internal image: possible regulation of a germ-line
A;Reference number: A91028; MUID:86136012; PMID:3937730
A;Accession: B28840
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B28840
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A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 1-98 <TIL>
A;Experimental source: B cell, strain [NZB x NZW]F1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin homology <TMM>
                                                                                                                                                                                                           A; Molecule type: mRNA
A; Residues: 1-101 <OLL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;10-84/Domain: immunoglobulin homology <IMM>
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C28840
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A; Residues: 1-101 <OLL>
                                                                          Matches
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                                                                                                              Query Match
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Best Local
                                                                                            Local
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83
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                                    1 LOYAIFPYT 9
                                                                  71.4%;
Similarity 77.8%;
7; Conservation
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LQYASSPYT
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91
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                                                                      pred. No. 7.6;
0; Mismatches
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Pred. No. 7.6;
0; Mismatches
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Pred. No. 7.4;
0; Mismatches
                                                                                            Score 35; I
Pred. No. 7
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RESULT 10
C71857
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A;Map position: 2p12-2p12
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer
C;Keywords: heterotetramer
E;16-90/Domain: immunoglobulin homology <IMM>
probable lipopolysaccharide biosynthesis protein - Helicobacter pylori (strain J99)
C;Species: Helicobacter pylori
A;Variety: strain J99
C;Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_change 09-Jul-2004
C;Accession: C71857
R;Alm, R.A.; Ling, L.S.L.; Moir, D.T.; King, B.L.; Brown, E.D.; Doig, P.C.; Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Goni, F.; Frangione, B.
Proc. Natl. Acad. Sci. U.S.A. 80, 48
A;Title. Namino acid sequence of the
A;Reference number: A9364; MUID:832
A;Accession: A01876
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A; Residues: 1-106 <SHL>
C; Superfamily: immunoglo
C; Keywords: heterotetra
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A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somati. A;Reference number: PL0231; MUID:90111618; PMID:2104919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:P01610 C;Comment: This chain was obtained
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;16-90/Domain: immunoglobulin homology <IMM>
;24-34/Region: complementarity-determining 1
;35-49/Region: framework 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Homo sapiens (man)
Date: 03-Aug-1984 #sequence_revision 03-Aug-1984 #text_change
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Keywords: heterotetramer; immunoglobulin
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Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
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e of the Fv region of a human monoclonal IgM
MUID:83273707; PMID:6410398
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Pred. No. 8.
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                        Query Match
Best Local S
Matches 4
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Peterson, S.; Loftus, B.; Richarusuu, Peterson, S.; Loftus, B.; Richarusuu, Peterson, S.; Loftus, B.; Richarusuu, Peterson, S.; Loftus, M.D.; Weidman, J.M.; Fujii, C.; Evennum, Son, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Smith, H.O.; Fraser, Nature 388, S39-547, 1997

A.Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A.; Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.

A.; Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.

A.; Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.
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submitted to the EMBL Data
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A; Residues: 1-398 <ARN>
A; Cross-references: UNIPROT: Q9ZKA7;
A; Cross-references: Strain J99
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A;Title: Genomic sequence comparison of two unrelated isolates of the human gastri A;Reference number: A71800; MUID:99120557; PMID:9923682
A;Accession: C71857
                                                                                              A;Molecule type: DNA
A;Residues: 1-431 <TOM>
A;Cross-references: UNIPROT:025733; GB:AE000617; GB:AE000511; NID:g2314256; PIDN:AAD08151
                                                                                                                                                                                                                                                                                                                                                                                                                           LPS biosynthesis protein - Helicobacter pylori (strain 26695)
C;Species: Helicobacter pylori
C;Date: 09-Aug-1997 #sequence_revision 09-Aug-1997 #text_change 09-Jul-2004
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A;Molecule type: DNA
A;Residues: 1-416 <ZIM>
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                                                                                                                                                                                                                                                                                                                            R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R.D.; Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKenneyson, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey, L.
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A;Molecule type: DNA
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Similarity 50.0%;
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55.6%;
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                       Score 34;
Pred. No.
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No. 47;
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Mismatches

Indels

Gaps

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Ig kappa chain V region - mouse (C;Species: Mus musculus (house mouse) (C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000 (C;Accession: S42466 R;Shiyanov, P.A.; Bespalov, I.A.; Terletskaya, H.N.; Deyev, S.M. submitted to the EMBL Data Library, March 1994 A;Reference number: S42466 A;Accession: S42466 A;Status: preliminary
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H84019
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F83707
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A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and A;Reference number: A83650; MUID:20512582; PMID:11058132
A; Molecule type: mRNA
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A;Experimental source: strain C-125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-107 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Takami, H.; Nakasone, K.; Takaki, Y.; Maeno, G.; Sasaki, R.; Masui, N.; Fuji, F.; Hira Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Recession: F83707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein BH0462 [imported] - Bacillus halodurans (strain C-125) C;Species: Bacillus halodurans C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004 C;Accession: F83707
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A;Molecule type: DNA
A;Residues: 1-97 <STO>
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Thes 5; Conserva
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A;Cross-references: EMBL:X78108; NID:g460824; PIDN:CAA54998.1; PID:g460825
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;26-100/Domain: immunoglobulin homology <IMM>
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Search completed: April Job time : 5.2492 secs
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                                                           100 QYSSYPYT 107
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49
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81.6
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l: uniprot_sprot:*
2: uniprot_trembl:*
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Copyright (c) 1993 - 2005
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Q9VIC3
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Q98UD2
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Q96y99
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Q96189
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Q7g908
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Q92169
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           8 anopheles g
4 drosophila
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Q96YG9;
01-DEC-2001
01-DEC-2001
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SEQUENCE
 SEQUENCE FROM N.A.
                     NCBI_TaxID=57060;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sulfolobus tokodaii.
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6; Conserve
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67.3	67.3	69.4	69.4	69.4	69.4	69.4	69.4	69.4	69.4	71.4	71.4	71.4	71.4
97	79	870	779	431	425	416	398	348	325	2396	945	879	879
N	N	N	N	N	N	N	N	N	N	N	N	Ŋ	N
Q9К8Р3	Q8GR38	Q815P3	Q7S947	025733	Q82VZ6	Q9P329	Q9ZKA7	Q6WN37	Q6LQ11	Q7UI32	Q7R3B3	Q9VDC6	076858
Q9k8	Q8gr	Q8i5p3	Q78947	025733	Q82vz6	Q9p329	Q9zka7	Q6wn37	Q61q1:	Q7u132	Q7r3b3	Q9vdc6	076858
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bacillus ha	enterococcu	plasmodium	neurospora	helicobacte	nitrosomona	schi zosacch	helicobacte	eleutheroda	photobacter	rhodopirell	giardia lam	drosophila	drosophila

ALIGNMENTS

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Q7ZZM4 PRELIMINARY;
Q7ZZM4;
01-JUN-2003 (TrEMBLrel. 2
01-JUN-2003 (TrEMBLrel. 2
01-MAR-2004 (TrEMBLrel. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.

STRAIN-JCM 10545 / 7;

MEDLINE-21456156; PubMed=11572479;

MEDLINE-21456156; PubMed=11572479;

Kawarabayasi Y., Hino Y., Horikawa H., Jin-no K., Takahashi M.,

Sekine M., Baba S.-I., Ankai A., Kosugi H., Hosoyama A., Fukui S.,

Nagai Y., Nishijima K., Otsuka R., Nakazawa H., Takamiya M., Kato Y.,

Yoshizawa T., Tanaka T., Kudoh Y., Yamazaki J., Kushida N., Oguchi A.,

Aoki K.-I., Masuda S., Yanagii M., Nishimura M., Yamagishi A.,

Oshima T., Kikuchi H.;

Complete genome sequence of an aerobic thermoacidophilic

Crenarchaeon, Sulfolobus tokodaii strain7.";
       VegT.
Eleutherodactylus coqui (Puerto Rican coqui).
Eleutherodactylus coqui (Puerto Rican coqui).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-DEC-2001 (TrEMBLrel. 19, 01-DEC-2001 (TrEMBLrel. 19, 01-JUN-2003 (TrEMBLrel. 24, Hypothetical protein ST2202, OrderedLocusNames=ST2202;
Eukaryota; Metazoa; Chordata; Cra
Amphibia; Batrachia; Anura; Neoba
Telmatobiinae; Eleutherodactylus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA Res. 8:123-140(2001).
EMBL; AP000989; BAB67308.1; -
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173 AA; 19383 MW; 14F1F8AAL
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26,
                                                                                                                           Last sequence update)
Last annotation update)
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Pred. No. 5.4;
3; Mismatches
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Last annotation update)
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                                                    Euteleostomi;
                          Leptodactylidae;
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Q654C)
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AC Q654
DT 25-C
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                                         D 029288 PRELIMINAKY; ...,
C 029288; PRELIMINAKY; ...,
T 01-JAN-1998 (TrEMBLrel. 05, Created)
T 01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
JT 01-JUN-2003 (TrEMBLrel. 24, Last amotation update)
B Bile acid-inducible operon protein F (BaiF-2).
GN OrderedLocusNames=AF0974;
OS Archaeoglobus fulgidus.
OC Archaea; Euryarchaeota; Archaeoglobi; Archaeoglobales
OC Archaeoglobaceae; Archaeoglobus.
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PRINTS; PR00937; TBOX.
SMART; SM00425; TBOX; 1.
PROSITE; PS01283; TBOX 1; 1.
PROSITE; PS50252; TBOX 3; 1.
SEQUENCE 417 AA; 47815 MW;
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Gukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

Ehrhartoideae; Oryzeae; Oryza.

NCBI_TaxID=39947;
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Q654Q7;
       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AP004651; BAD45710.1; -.
Hypothetical protein.
SEQUENCE 135 AA; 14500 MW; C2A82C97536D420B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sasaki T., Matsumoto T., Yamamoto "Oryza sativa nipponbare(GA3) gencolone:OSJNBa0091006.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
4 Hypothetical protein OSJNBa0091G06.4.
Name=OSJNBa0091G06.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GO; GO:0005634; C:nucleus; IEA.
GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:0006355; P:regulation of transcription, DNA-dependent;
InterPro; IPR008967; P53 like DNA_bnd.
InterPro; IPR001699; TF_T-box.
Pfam; PF00907; T-box; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
HSSP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beckham Y.M., Nath K.,
Submitted (MAR-2003) to
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%L; AY251031; AAP20869.1; -.

P; O15119; 1H6F.

GO:0005634.
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YAVFPYT 123
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85.7%;
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75.0%;
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Pred. No. 10;
1; Mismatches
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Pred. No. 20;
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RESULT 5
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ID Q7RXX3
AC Q7RXX
AC Q7RXX
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Affe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,
Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,
RA Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,
RA Cui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,
RA Cui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,
RA Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,
ROY A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S.,
RA Kamal M., Kamvysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,
RA Kamal M., Kamvysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,
RA Katystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,
ROGODNI C., Macino G., Catcheside D., Li W., Pratt R.J., Osmani S.A.,
RA Krystofova S., Glass L., Orbach M.J., Berglund J., Voelker R.,
RA Varden O., Plamann M., Seiler S., Dunlap J., Radford A., Aramayo R.,
RA Matvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,
RA Matvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,
RA Paulsen I., Sachs M.S., Lander E.S., Nusbaum C., Birren B.,
"The Genome Sequence of the Filamentous Fungus Neurospora crassa.";
Nature 0:0-0(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
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Q7RXK3;
01-MAR-2004
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Fungi, Ascomycota; Pezizomycotina; Sordariomycetes; Sordariomycetidae; Sordariales; Sordariaceae; Neurospora. NCBI_TaxID=5141;
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SEQUENCE
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01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation updat
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Nature 390:364-370(1997).
EMBL; AE001037; AAB90269.1; -.
PDR; F69371, F69371.
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MEDLINE=98049343; PubMed=9389475; DOI=10.1038/37052;

Klenk H.-P., Clayton R.A., Tomb J.-F., White O., Nelson K.E.,

Ketchum K.A., Dodson R.J., Gwinn M.L., Hickey E.K., Peterson J.D.,

Richardson D.L., Kerlavage A.R., Graham D.E., Kyrpides N.C.,

Fleischmann R.D., Quackenbush J., Lee N.H., Sutton G.G., Gill S.R.,

Fleischmann R.D., Quackenbush J., Lee N.H., Sutton G.G., Gill S.R.,

Kirkness E.F., Dougherty B.A., McKenney K., Adams M.D., Loftus B.J.,

Peterson S.N., Reich C.I., McNeil L.K., Badger J.H., Glodek A.,

Zhou L., Overbeek R., Gocayne J.D., Weidman J.F., McDonald L.A.,

Utterback T.R., Cotton M.D., Spriggs T., Artiach P., Kaine B.P.,

Sykes S.M., Sadow P.W., D'Andrea K.P., Bowman C., Fujii C.,

Garland S.A., Mason T.M., Olsen G.J., Fraser C.M., Smith H.O.,

Whoese C.R., Venter J.C.;
                                                                                                                                  -!- SIMILARITY: Belongs to the type-B carboxylesterase/lipase family.
-!- CAUTION: The sequence shown here is derived from an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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TIGR; AF0974; -.
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                              preliminary data.
L; AABX01000743; E
                                                                                                EMBL/GenBank/DDBJ whole
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                              EAA27327.1;
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Pred. No. 34;
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RESULT 7
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Best Local S
Matches 5
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Rd Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Read N.D.,
Rd Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Read N.D.,
AA Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,
Rd Rikins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,
Rd Qui D., Ianakiev P., Pedersen D., Nelson M., Waabburne M.,
Rd Qui D., Ianakiev P., Pedersen D., Nelson M., Waabburne M.,
Rd Qui D., Ianakiev P., Pedersen D., Nelson M., Waabburne M.,
Rd Qui D., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,
Rd Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,
Rd Kothe G.O., Jedd G., Mewes W., Staben C., Marcotte E., Greenberg D.,
Rd Roy A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S.,
Ramal M., Kamvyssells M., Maucell E., Bielke C., Rudd S., Frishman D.,
Rcyatofova S., Rasmussen C., Mezzenberg R.L., Perkins D.D., Kroken S.,
Cogoni C., Macino G., Catcheside D., Li W., Pratt R.J., Osmani S.A.,
DeSouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R.,
Varden O., Planann M., Seller S., Dunlap J., Radford A., Aramayo R.,
Natvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,
Paulsen I., Sachs M.S., Lander E.S., Nusbaum C., Birren B.;
"The Genome Sequence of the Filamentous Fungus Neurospora crassa.";
Nature O:O-O(2003).
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Q75014
01-MAR-2004 (TrEMBLrel 2:
01-MAR-2004 (TrEMBLrel 2:
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HSSP; P22303; 1F8U.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preliminary data.
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MEDLINE=96337999; PubMed=8688087;

Bult C.J., White O., Oleen G.J., Zhou L., Fleischmann R.D.,

Bult C.J., White O., Oleen G.J., Zhou L., Fleischmann R.D.,

Sutton G.G., Blake J.A., FitzGerald L.M., Clayton R.A., Gocayne J.D.,

Korlavage A.R., Dougherty B.A., Tomb J.-F., Adams M.D., Reich C.I.,

Overbeek R., Kirkness E.F., Weinstock K.G., Merrick J.M., Glodek A.,

Scott J.L., Geoghagen N.S.M., Weidman J.F., Fuhrmann J.L., Nguyen D.,

Utterback T.R., Kelley J.M., Peterson J.D., Sadow P.M., Hanna M.C.,

Cotton M.D., Roberte K.M., Hurst M.A., Kaine B.P., Borodovsky M.,

Klenk H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.,

"Complete genome sequence of the methanogenic archaeon, Methanococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q57895;
01-NOV-1997
01-NOV-1997
05-JUL-2004
            SEQUENCE FROM N.A.

STRAIN-CGA009 / ATCC BAA-98;
PubMed=14704707; DOI=10.1038/nbt923;
Larimer F.W., Chain P., Hauser L., Lamerdin J.E., Malfatti S.,
Land M.L., Pelletier D.A., Beatty J.T., Lang A.S., Tabita F.R.,
Gibson J.L., Hanson T.E., Bobst C., Torres y Torres J.L., Peres
Harrison F.H., Gibson J., Harwood C.S.;
"Complete genome sequence of the metabolically versatile
photosynthetic bacterium Rhodopseudomonas palustris.";
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SEQUENCE
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                                                                                                                                                                                                            Rhodopseudomonas pallustria.
Bacteria, Proteobacteria, Alphaproteobacteria,
Bradyrhizobiaceae, Rhodopseudomonas.
                                                                                                                                                                                                                                                             05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Putative Affr-binding protein of ABC transporter.
OrderedLocusNames=RPA0468;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Archaea; Euryarchaeota; Methanococci; Methanococcales;
Methanocaldococcaceae; Methanocaldococcus.
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C Probably responsible for the translocation of the substrate across
C the membrane (By similarity).
C -!- SUBCELLULAR LOCATION: Integral membrane protein (By similarity).
C -!- SIMILARITY: Belongs to the binding-protein-dependent transport
C SYSTER BELLY K5725912; j. -.
C EMBL; MX572594; CAR25912; j. -.
C GO: 0016021; C:integral to membrane; IEA.
C GO: GO:0005524; F:ATP binding; IEA.
C GO: GO:0005215; F:transporter activity; IEA.
C GO: GO:0005215; F:transporter activity; IEA.
C GO: 00005215; BPD_transporter activity; IEA.
C GO:0005215; BPD_transporter activity; IEA.
C GO:0005216; BPD_transporter activity; IEA.
C GO:0005
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Q7P6U2;
Q1P6U2;
Q1-MAR-2004 (TrEMBLrel. 2
Q1-MAR-2004 (TrEMBLrel. 2
Q1-MAR-2004 (TrEMBLrel. 2
                                                             Q87HF9 PRELIMINARY; PRT; 30 Q87HF9; (TrEMBLrel. 24, Created) 01-JUN-2003 (TrEMBLrel. 24, Last sequen 01-MAR-2004 (TrEMBLrel. 26, Last annota Transcriptional regulator, LysR family. CreeredLocusNammes-VPA1006;
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EMBL; AABF01000027; E:
Hypothetical protein.
SEQUENCE 276 AA; 3:
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                                 Vibrio parahaemolyticus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       49256;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    φ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32878 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EAA24518.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73.5%;
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Fusobacterales; !
      Gammaproteobacteria; Vibrionales;
                                                                                                                                    Last sequence update)
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Pred. No. 81;
3; Mismatches
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Pred. No.
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Last annotation updat
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Best Local S
Matches 5
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Best Local &
                                                                                                      PROSITE;
Complete
SEQUENCE
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STRAIN-RIMD 2210633 / Serotype O3:K6;

MEDLINE=22508454; PubMed=12620739; DOI=10.1016/S0140-6736(03)12659-1;

Makino K., Oshima K., Kurokawa K., Yokoyama K., Uda T., Tagomori K.,

Iijima Y., Najima M., Nakano M., Yamashita A., Kubota Y., Kimura S.,

Yasunaga T., Honda T., Shinagawa H., Hattori M., Iida T.;

"Genome sequence of Vibrio parahaemolyticus: a pathogenic mechanism
                                                                                                                                                                                                                                                                                                           high pressure adaptations.";
Submitted (MAR-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; CR3708670; CAG20746.1; -.
GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:0006355; P:regulation of transcription, DNA-dependent; IEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OGLENO;
OS-JUL-2004 (TrEMBLrel. 27, Created)
OS-JUL-2004 (TrEMBLrel. 27, Last sequence update)
OS-JUL-2004 (TrEMBLrel. 27, Last annotation update)
OS-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical transcriptional regulator, LysR family.
Name=VV21654; OrderedLocusNames=PBPRA2361;
Photobacterium profundum (Photobacterium sp. (strain SS9)).
Bacteria; Proteobacteria; Gammaproteobacteria; Vibrionales;
Vibrionaceae; Photobacterium.
                                                                                                                                                InterPro; IPR000847; HTH LysR.
InterPro; IPR005119; LysR subst.
InterPro; IPR009058; Wing_hlx_DNA_bnd.
Pfam; PF00126; HTH 1; 1.
Pfam; PF03466; LysR substrate; 1.
PROSITE; PS0931; HTH_LYSR; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000847; HTH_LysR.
InterPro; IPR005119; LysR subst.
InterPro; IPR009058; Wing_hlx_DNA_bnd.
Pfam; PF00126; HTH 1; 1.
Pfam; PF03466; LysR substrate; 1.
PFINTS; PR00039; HTHLYSR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vezzi A., Campanaro S., D
Cestaro A., Malacrida G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q6LPN0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -1- SIMILARITY: Contains 1 HTH lysR-type DNA-binding domain.
EMBL; AP005087; BACC2349.1; -.
GO; GO:0003700; Firranscription factor activity; IEA.
GO; GO:0006355; Piregulation of transcription, DNA-dependent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               distinct from that of V. cholerae."; Lancet 361:743-749(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=74109;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Valle G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Genome analysis of Photobacterium
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Similarity 5; Conserv
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301 AA;
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                      73.5%;
                                                                                                      33862 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          D'Angelo M., Simonato F., Vitulo N., . Simionati B., Cannata N., Bartlett
  2
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                        Score 36; DB
Pred. No. 89;
                                                                                                      3A6D90E214EDA8E0
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  Mismatches
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GN Names
OS Xenog
OC Amphla
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RL SLUBAT
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Best Local
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YEST XENIA

STANDARD; PRT; 455 AA.

P87377; P79930; P87386;

P87377; P79930; P87386;

P87377; P79930; P87386;

P87377; P79930; P87386;

P1-NOV-1997 (Rel. 35, Last sequence update)

O1-NOV-1997 (Rel. 35, Last sequence update)

O5-JUL-2004 (Rel. 44, Last amotation update)

T-box protein VEGT (T-box protein BRAT) (T-box protein antipodean)

Name=VEGT; Synonyms=APOD, BRAT;

Xenopus laevis (African clawed frog).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelecstomi;

Eukaryota; Metazoa; Chordata; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUN-2001
01-JUN-2001
01-MAR-2004
Zhang J., King M.L.; "Xenopus VegT RNA is localized to the vegetal cortex during oogenesis "Xenopus VegT RNA is localized to the vegetal cortex during oogenesis and encodes a novel T-box transcription factor involved in mesodermal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HSSP; 015119; 1H6P.

GO; GO:0005634; C:nucleus; IEA.
GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:0006355; P:regulation of transcription, DNA-dependent;
InterPro; IPR008967; P53_like_DNA_bnd.
InterPro; IPR001699; TF_T-box.
Pfam; PF00907; T-box; 1.
PRINTS; PR00327; TBOX; 1.
SMART; SM00425; TBOX; 1.
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Bubunenko M., Vempati U.D.,
Submitted (AUG-1999) to the
EMBL; AF180351, AAK00596.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xenopus borealis (Kenyan clawed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                 Horb M.E.,
                                                                                                                                                                                                                                                                              TISSUE-Embryo;
                                                                                                                                                                                                                                                                                                                                                                                  Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q98UD2,
                                                                                                 TISSUE=Oocyte;
MEDLINE=97164724;    PubMed=9012531;
                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE; PS01283; TBOX_1; 1.
PROSITE; PS01284; TBOX_2; 1.
PROSITE; PS50252; TBOX_3; 1.
SEQUENCE 454 AA; 51855 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=VegT;
                                                                                                                                                                 SEQUENCE FROM N.A.
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YAVYPYT 276
                                                                                                                                                                                                                       , Thomsen G.H.; (MAR-1997) to
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4 (TrEMBLrel. 26, 1
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                                                                                                                                                                                                                    the
                                                                                                                                                                                                                    EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        King M.L.;
EMBL/GenBank/DDBJ databases
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Last annotation update)
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 Mismatches

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Pred. No. 1.3e+02;
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RESULT 14
013161
ID 01316
AC 01316
DT 01-JU
DT 01-JU
DT 25-OC
DE T-box
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                                                                                                                                                                                                                                                                                          Query Match
Ol3161 PRELIMINARY; PKT; 455 Am.
Ol3161;
Ol3101;
Ol-JUL-1997 (TrEMBLrel. 04, Created)
Ol-JUL-1997 (TrEMBLrel. 04, Last sequence update)
25-CCT-2004 (TrEMBLrel. 28, Last annotation updat)
T-box protein (Xombi protein).
                                                                                                                                                                                                                                                                                                                              CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stennard F., Carnac G., Gurdon J.B.;

"The Xenopus T-box gene, Antipodean, encodes a vegetally localised maternal mRNA and can trigger mesoderm formation.";

Development 12:4179-4188(1996).

-i- FUNCTION: Transcription factor involved in mesodermal patterning. Appears to pattern mesoderm along the dorsoventral and posterior axis. It activates WNT-8, EOMES and brachyury expression.

-i- SUBCELLULAR LOCATION: Nuclear.

-i- TISSUE SPECIFICITY: Expressed both maternally and zygotically. Maternally localized to the egg vegetal hemisphere, and in the developing embryo, in the posterior paraxial mesoderm and ventral blastopore.
                                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PR00937; TBOX.

PROSITE; PS01283; TBOX.2; 1.

PROSITE; PS01264; TBOX.2; 1.

PROSITE; PS50252; TBOX.3; 1.

Developmental protein; DNA-binding; Nuclear protein;

Transcription regulation.

Transcription 730

DNA_BIND

57

230

T-box.
                                                                                                                                                                                                                                                                                                                                                                                                Transcription 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR008967; P53 like DNA bnd InterPro; IPR001699; TF T-box. Pfam; PF00907; T-box; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMMBL, U89707; AAB49478.1; -.
EMMBL, U59483; AAB93301.1; -.
EMMBL, X99905; CAA68179.1; -.
HSSP; O15119; 1H6F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation between the Swiss Institute of Bioinformatics are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ----
                                                                                                                                                                                                                                                                                                                                                                                CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REVISIONS TO 209 AND 396-455.

Zhang J., King M.L.;
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   patterning.";
Development 122:4119-4129(1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=97164730; PubMed=9012537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INDUCTION: By TGF-beta family members.
SIMILARITY: Contains 1 T-box domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DEVELOPMENTAL STAGE: Maternally expressed from early oogenesis. Zygotic expression occurs from late blastula and reaches maximum levels during gastrulation (stages 10.5-12). Levels decline at the time of blastopore closure (stage 13).
                                                                                                                                                                                           446
                                                                                                                                                                                                                        2 QYAIFPYT 9
                                                                                                                                                                                                                                                          Similarity
5; Conserv
                                                                                                                                                                                        QYSLFPYS
                                                                                                                                                                                                                                                                                                                              244
455 AA;
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                                                                                                                                                                                           453
                                                                                                                                                                                                                                                                                                                                                 244
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25
                                                                                                                                                                                                                                                                                                                                51795 MW;
                                                                                                                                                                                                                                                                            73.5%;
62.5%;
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                                                                                                                                                                                                                                                                            Score 36; DB 1;
Pred. No. 1.3e+02;
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H -> L (in Ref. 2 and 3).
                                                                                                                                                                                                                                                                                                                                9DD12CD704F2AE07 CRC64;
                                   update)
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Name=Xombi; Xenopus laevis (African clawed frog). Xenopus laevis (African clawed frog). Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleosto Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;

(African clawed frog). nzoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Xenopodinae; Xenopus.

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SON SON REPRESENTATION OF SON REPRESENTATION
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Branstein M.J., Ugdin T.B., Toshiyuki S., Caraninci P., Prange C.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Caraninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robask S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Willalon D.K., Mozny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Hahtesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Kozrywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.:
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Best Local S
Matches 5
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                                                                                                                                                                                                             GO; GO:0005634; C:nucleus; IEA.
GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:000355; F:regulation of transcription, DNA-dependent; IEA.
GO; GO:0006355; F:regulation of transcription, DNA-dependent; IEA.
InterPro; IPR008967; P53 like DNA_bnd.
InterPro; IPR001699; TF_T-box.
Pfam; PF00907; T-box; 1.
PRINTS; PR00937; TBOX; 1.
PROSITE; PR00317; TBOX; 1.
PROSITE; PS01264; TBOX 1; 1.
PROSITE; PS01264; TBOX 3; 1.
PROSITE; PS01264; TBOX 3; 1.
SEQUENCE 455 AA; 51963 MW; 2B9C8FBCCD78E37D CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Oocytes;
Klein S., Strausberg
Submitted (MAY-2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22341132; PubMed=12454917; I
Klein S.L., Strausberg R.L., Wagner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Genetic and genomic tools for Xenopus research: The NIH Xenopus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Oocytes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            S., Strausberg R.;
tted (MAY-2004) to the EMBL/GenBank/DDBJ
S83518; AAB50917.1; -.
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                                                                                         Similarity
5; Conserv
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                                                                                         Score 36; DB 2;
Pred. No. 1.3e+02;
3; Mismatches 0
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r L., Pontius J., Clifton S.W.,
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                                                                                                   0;
                                                                                                                                                               Length 455;
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Matches 5
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SMART; SM00425; TBOX; 1.
PROSITE; PS01283; TBOX 1; 1.
PROSITE; PS01284; TBOX 2; 1.
PROSITE; PS50252; TBOX 3; 1.
SEQUENCE 455 AA; 51890 MW;
  time
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01-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. Bubunenko M., Vempati U.D., Submitted (AUG-1999) to the EMBL; AF180352; AAK00597.1; HSSP; O15119; 1H6F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xenopus tropicalis (Western clawed frog) (
Eukaryota; Metazoa; Chordata; Craniata; Ve
Amphibia; Batrachia; Anura; Mesobatrachia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q98UD1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GO; GO:0005634; C:nucleus; IEA.
GO; GO:0003700; F:transcription factor
GO; GO:0006355; P:regulation of transc
GO; GO:006355; P:regulation of transc
InterPro; IPR0018967; P53 like DNA_bnd.
InterPro; IPR001699; TF_T-box.
Pfam; PF00907; T-box; 1.
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01-MAR-2004
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  completed: April
me : 20.2109 secs
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5; Conserv
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|QYSLFPYS
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                                                                                                                 453
                               4,
                                                                                                                                                                                                                                                73.5%;
                                  2005, 15:56:19
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17,
26,
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EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                          Score 36; DB 2;
Pred. No. 1.3e+02;
3; Mismatches (
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                                                                                                                                                                                                                                                                                                                                     0D2502AF91CBA2B0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              factor activity; IEA.
transcription, DNA-dependent; IEA
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ia; Pipoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Silurana tropicalis)
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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1: geneseqp1980s:*
2: geneseqp1990s:*
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193.690 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Regult No.	Score	Query	Length	B	ID	Description
, 	30	88	1 U1	N	AAR85496	
2	30	100.0	ហ	N	AAW24504	Aaw24504 Mouse
w	30	00.	Մ	w	AAY32223	
4	30	00.	ហ	σ	AAU80973	ω
տ	30		տ	7	ABR61517	7
6	30	•	ري ري	8	ADG34365	
7	30	100.0	₅	8	ADL23057	Adl23057 Myelin
8	30	•	₅	8	ADS87336	
9	30	٠	S	8	ADS88789	_
10	30	•	6	4	AAB99237	_
11	30	•	7	N	AAY17489	Aay17489 Mab
12	30	100.0	7	ນ	AAY08698	
13	30	100.0	7	w	AAY83641	
14	30	•	10	N	AAW70611	۲
15	30	•	10	ហ	AAG80311	Aag80311 Anti-huma
16	30	٠	10	ហ	ABP61180	0
17	30	•	10	ထ	ADR89785	
18	30	٠	10	œ	ADS94361	-
19	30	100.0	12	7	ADC27954	
20	30	100.0	66	տ	ABP11252	
21	30	00.	67	7	ADH62639	Adh62639 Mouse
22	30	100.0	70	7	ADH62638	æ
23	30	8	70	7	ADH62637	-
24	30	100.0	74	ហ	ABP38658	_
у л	30	2				

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26
30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30	30
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0		100.0	100.0	100.0	100.0					100.0	100.0	100.0
116	116	116	116	116	116	116	116	116	116	116	116	116	116	115	115	112	112	92	92
ທ	u	5	5	ហ	ហ	ហ	ហ	ហ	ហ	σ	ហ	5	4	4	W	ŋ	N	Ŋ	N
ADG67533	ADG67531	ADG67543	ADG67537	ADG67541	AAE27834	AAE27832	AAE27828	AAE27826	AAE27825	AAE27837	AAE27839	AAE27830	AAB84739	AAB84740	AAY90823	AAU80980	AAW24521	AAU80997	AAW24529
Adg67533	Adg67531	Adg67543	Adg67537	Adg67541	Aae27834	Aae27832	Aae27828	Aae27826	Aae27825	Aae27837	Aae27839	Aae27830	Aab84739	Aab84740	Aay90823	Aau80980	Aaw24521	Aau80997	Aaw24529
KS antibo	Mouse KS	Variable	Variable	520C9 hyb	Human ant	Anti-Fact	Human ant	Anti-Fact											

ALIGNMENTS

AAR85496 standard; peptide;

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ARESULT 1
AARBAAAB
AN AARB
AX AARB
AX AARB
AX AARB
AX AARB
AX Scrv
XX Sing
XW tumo
CX Comp
CC 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SCFV (FRP) 5 CDR1H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAR-1996 (first entry)
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New bifunctional proteins for use in killing tumour cells - contg. a tumour antigen binding domain, a hinge region and a zeta chain derived from a T-cell antigen receptor.

WPI; 1995-393085/50.

Groner B,

Moritz D;

Disclosure; Page 6; 46pp; English.

Complementarity determining regions (CDRs) CDR1H, CDR2H and CDR3H (AAR85496-98) and CDR1L, CDR2L and CDR3L (AAR85502-04) are components of the heavy chain variable region and light chain variable region, respectively, of single chain antibody scFv(FRP)5 (AAR85494). The CDRs are derived from mouse monoclonal antibody FRP5 (ECACC 90112115). The scFv is specific for the extracellular domain of erbB-2, and is used as the tumour antigen binding domain of a bifunctional protein that is expressed in cytotoxic T-lymphocytes (CTLs), allowing adoptive immunotherapy of cancer

Sequence 5 AA;

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RESULT 2
AAW24504
ID AAW2
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 Query Match
Best Local S
Matches 5
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Best Local :
                                                                                            This peptide comprises complementarity determining region 1 (CDR1) of the heavy chain variable region (VH) of mouse anti-human Factor IX monoclonal antibody BC2 (see also ARW24519. Claimed humanised antibodies (see ARW24510-18) contain CDRs (see ARW24504-09) of BC2 VH and VI inserted into framework regions of selected human antibody sequences. They have self-limiting neutralising activity, and are useful as anticoagulant agents in treatment of thrombosis associated with myocardial infarct, unstable angina, atrial fibrillation, stroke, renal damage, pulmonary embolism, deep vein thrombosis, percutaneous transluminal coronary angioplasty, disseminated intravascular coagulation, sepsis, or artificial organs, shunts or prostheses (claimed). Also claimed are chimeric cantibodies, as well as Fab and F(ab')2 fragments. The claimed antibodies do not cause uncontrolled bleeding (contrast heparin and warfarin) since they provide only partial inhibition of coagulation
                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                             Claim 21; Page.65;
                                                                                                                                                                                                                                                                                                                                                                                            Inhibiting thrombosis with self-limiting antibody to coagulation factor avoids uncontrolled bleeding by providing only partial inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-385117/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Blackburn MN, Church WR, Feuers
Padlan EA, Patel AH, Sylvester
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-JAN-1996;
24-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    humanised antibody; antibody engineering; heavy chain; CDR; complementarity determining region; myocardial infarction; angiatrial fibrillation; stroke; kidney damage; pulmonary embolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9726010-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SMIK ) SMITHKLINE BEECHAM CORP.
(UYVE-) UNIV VERMONT & STATE AGRIC COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disseminated intravascular coagulation; artificial organ; sepsis; shunt;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thrombosis; therapy; Factor IX; anticoagulant; monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse anti-human Factor IX antibody BC2 heavy chain CDR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          deep vein thrombosis; coronary angioplasty;
                 Local Similarity
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 100.0%; Score 30; DB 2; ilarity 100.0%; Pred. No. 1.8e+06; Conservative 0; Mismatches 0;
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96US-0029119P.
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                                                                                                                                                                                                                                                                                                                                                             150pp; English
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100.0%; Pred. No. 1.8e+06;
. Mismatches 0;
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                                 Length 5;
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RESULT 3
ID AAY3
ID AAY3
XX AC AAY3
XX G-CS
XX COME
XX MUS
XX MI

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AAU80973
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents complementarity determining region 1 (CDR1) of the heavy chain variable region of murine monoclonal antibody mabl63-93. This antibody is an example of an agonist molecule that specifically binds to or interacts with human granulocyte colony stimulating factor (G-CSF) receptor to stimulate cell proliferation and differentiation, especially by dimerising the receptor or activating phosphorylation of kinases associated with the receptor. Agonist antibodies can be used to stimulate proliferation of G-CSF-dependent cells, e.g. to differentiate leading to a repopulation of neutrophilic granulocyte lineage cells, especially to treat neutropenia (claimed). They can also be used to detect human G-CSF receptor immunologically
                    Murine BC2 heavy chain variable region CDR1.
                                                                                09-APR-2002
                                                                                                                                         AAU80973;
                                                                                                                                                                                            AAU80973 standard; peptide; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neutropenia by stimulating proliferation of neutrophilic
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                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Pred. No. 1.8e+06;
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RESULT 5
ABR61517
ID ABR6
XX ABR6
XX ABR6
XX ABR6
XX LST
DT 15-J
XX Huma
XX EGP-
KW RS7
XX HOMC
XX HOM
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; mouse; BC2; animal post-thromboembolic induced ischaemia; thrombolytic agent; anti-factor IX antibody; plasminogen activator; thromboembolic stroke; cerebroprotective; anti-cagqulant; Thrombolytic; vasotropic; cardiant; anti-respiratory syncytial virus;
                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Humanised RS7 antibody heavy chain variable region CDR1 fragment.
                                                   01-MAR-2002; 2002US-0360229P
                                                                                                                                    03-MAR-2003; 2003WO-GB000885
                                                                                                                                                                                                                                                                                                                   WO2003074566-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EGP-1; glycoprotein; cytostatic; antigen binding protein; cancer; RS7 antibody; complementarity determining region; CDR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating post-thromboembolic induced ischemia in an animal by administering anti-factor IX antibody in combination with a plasminogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-082944/11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAY-2000; 2000US-00571434.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200187339-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         method
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Š
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 30; DB 5;
Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Toomey JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 5;
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RESULT 6
ADG34365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an antibody (Ab) or fragment that binds to an CC EGP-1 glycoprotein. The methods are useful for delivering a diagnostic or therapeutic composition, diagnosing or treating cancer, treating a CC malignancy, and treating a cancer cell. A therapeutic conjugate or CC diagnostic conjugate comprising EGF-1 Mab, a fragment of an antibody CC fusion protein or fragment is used for the diagnosis or treatment of a CC malignancy when bound to at least one therapeutic agent. The diagnostic/ CC therapeutic agent or a combination of both comprising at least one Ab are CC used for targeting an EGP-1 target antigen on a cell. The cancers are CC used for targeting an EGP-1 target antigen on a cell. The cancers are CC used for targeting an EGP-1 target antigen on a cell. The cancer. Naked anti-EGP-1 Mab or fragment is used for the treatment of cancer. The CC invention provides multivalent, monospecific binding proteins that are useful in a direct targeting system and for production of multivalent, comparison of multivalent are useful in an affinity enhancement CC system. The present sequence represents a humanised anti-EGP-1 antibody constant variable region CDR1 fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                       04-DEC-2003
                                                                                                                                                                                                                                                                          da ap
                                                                                                                                                                                                                                                                                                        humanised monoclonal antibody; complementarity determining region; CDR; murine monoclonal antibody BAT-1; mBAT-1; framework region; FR; acceptor human immunoglobulin; anti-tumour; immunogenic; cytostatic; anti-HIV; immunotherapy; cancer; HIV infection; autoimmune disorder; immune deficiency; mouse; murine; heavy chain variable region.
              WPI; 2004-035027/03
                                                Hardy B,
                                                                                                                                    23-MAY-2002; 2002IL-00149820
                                                                                                                                                                    22-MAY-2003; 2003WO-IL000425
                                                                                                                                                                                                                                         WO2003099196-A2
                                                                                                                                                                                                                                                                                                                                                                                                                   Humanised
                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG34365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG34365 standard; peptide; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 6; Page 65; 97pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monovalent and multivalent humanized RS7 antigen binding proteins useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Govindan S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IMMU-) IMMUNOMEDICS INC. (MCCA/) MCCALL J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the diagnosis and therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              μ
                                                                                 CURE TECH LTD.
MOR RES APPL LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NYGMN 5
                                                Jones ST,
                                                                                                                                                                                                                                                                                                                                                                                                                 mouse BAT-1 antibody-derived HC variable region SeqID9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                Klapper L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 30; DB 7; Pred. No. 1.8e+06; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Goldenberg DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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RESULT 7
ADL23057
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Best Local :
New altered antibody that binds to and neutralizes myelin associated glycoprotein (MAG), useful for preparing a composition for treating or preventing stroke or other neurodegenerative disorders e.g., Alzheimer's
                                                                                            N-PSDB; ADL23034.
                                                                                                                                                       Ellis
                                                                                                                                                                                                                                  06-AUG-2002; 2002GB-00018229
06-AUG-2002; 2002GB-00018230
06-AUG-2002; 2002GB-00018230
06-AUG-2002; 2002GB-00018234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to a novel humanised monoclonal antibody having at least one complementarity determining region (CDR) of murine monoclonal antibody BAT-1 (mEAT-1) and a framework region (FR) derived from an acceptor human immunoglobulin. The humanised antibody retains the anti-tumour activity of mBAT-1 monoclonal antibody and is less immunogenic in a human subject than the murine antibody. The invention may be useful in the development of compounds with a cycostatic or anti-HIV activity. The human monoclonal antibodies are useful in immunotherapy, particularly for treating cancer. The antibodies are also useful for treating early stages of HIV infection or in patients whose blood count shows a decrease in autoimmune disorders, or in some cases of genetic or acquired immune
                                                                                                                                                                                                                                                                                                                                        05-AUG-2003; 2003WO-EP008749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Myelin associated glycoprotein MAG binding antibody heavy chain
                                                                                                                                                                                                                                                                                                                                                                                   19-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                          WO2004014953-A2
                                                                                                                                                                                                (GLAX )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neurodegenerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADL23057 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          deficiencies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acceptor human
ages of HIV.
                                                                                                              2004-180641/17.
                                                                                                                                                       Ä,
                                                                                                                                                                                                GLAXO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MAG; myelin associated
                                                                                                                                                     Germaschewski V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                              GROUP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorder;
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100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 glycoprotein; heavy chain; CDR; stroke; therapy; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.8e+06;
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믕 5

The present invention fragment, which binds

relates to to and neul

s to a new altered a neutralizes myelin

antibody or n associated

its functional glycoprotein

detecting

or treating cancer or ischemic

lesion.

Claim 1; Page

53; 67pp; English.

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RESULT 8
ADS87336
ID ADS88
XX ADS88
XX ADS88
XX Cyto
CYto
KW Cyto
KW Cyto
KW Cohim
KW Cinfil
KW 
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Best Local S
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                      that binds NCA90 or NCA95, where when the Mab or its fragment binds NCA90 the Mab or its fragment is chimeric, partially humanized or fully humanized and where when the Mab or its fragment binds NCA95 the Mab or its fragment is either fully humanized or chimeric, partially humanized or fully humanized BM 250/183. (I) is useful for treating, detecting or imaging sites of inflammation resulting from appendicitis, inflammatory bowel disease, pelvic inflammatory disease, fever and cystic fibrosis and treating granulocyte related disorders, atherosclerosis and infarction.
                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 105; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                  Monoclonal anti-granulocyte antibody that binds NCA90 and NCA95 useful for treating malignancy, inflammation, atherosclerosis, infarction, or other granulocyte related disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Goldenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiatherosclerotic; cardiant; monoclonal antibody; NCA90; NCA95; chimeric antibody; humanized antibody; inflammation; appendicitis; inflammatory bowel disease; pelvic inflammatory disease; fever; cystic fibrosis; granulocyte related disorder; atherosclerosis; infarction; cancer; ischemic lesion; complementarity determining r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-329873/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-SEP-2002; 2002US-0414341P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-SEP-2003; 2003WO-GB004229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; vasotropic; antimicrobial; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Humanised antibody MN3 heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADS87336
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MAG) and comprises a light chain variable domain (VL) comprising complementary determining region light 1 (CDRL1), CDRL2 or CDRL3 and/or a heavy chain variable domain (VH) comprising CDRH1, CDRL2 or CDRH3. The antibody is useful for preparing a composition for treating or preventing stroke or other neurodegenerative disorders in a human, e.g., traumatic brain injury, Alzheimer's disease, dementias, peripheral neuropathy, Parkinson's disease, Huntington's disease and multiple sclerosis. The present sequence is the anti-MAG antibody heavy chain CDR H1 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention.
                                                                                                                                                                                                                                                                                 invention relates to a monoclonal antibody (Mab)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MCCALL J
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Pred. No. 1.8e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                The specification describes a method for promoting oligodendrocyte survival in a human suffering or at risk of developing stroke or another neurological diseases. The method comprise administering to the human an anti-myelin-associated glycoprotein (MAG) antibody or its functional fragment. The anti-MAG antibody or its functional fragment is useful in the manufacture of a medicament for the promotion of oligodendrocyte
                                                                                                                                                                                                                                                                                         survival in a human suffering from or at risk of developing stroke or another neurological disease. They can also be used in treating neurological diseases, such as Alzheimer's diseases, multiple sclerosis and/or stroke, by promoting oligodendrocyte survival. ADS88789-ADS88791 represent the heavy chain complementarity determining regions (CDRs) of an anti-MAG antibody which can be used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            myelin-associated glycoprotein; MAG; anti-MAG antibody; Alzheimer's disease; multiple sclerosis; chain complementarity determining region; CDR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Promoting oligodendrocyte survival in humans with neurological diseases, such as Alzheimer's disease, multiple sclerosis and/or stroke, using an anti-myelin-associated glycoprotein (MAG) antibody.
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                CC chordoma, anglosarcoma, emidate and traing exclusions. CC chordoma, anglosarcoma, epithelial adenocarcinoma, and liver metastases CC thereof, lymphangiosarcoma, lymphangioendotheliosarcoma, hepatoma, CC cholangiocarcinoma, synovioma, mesothelioma, Ewing's tumour, CC rhabdomyosarcoma, colon carcinoma, basal cell carcinoma, sweat gland Ccarcinoma, papillary carcinoma, mesothelioma, Ewing's tumour, CC adenocarcinoma, cystadenocarcinoma, medullary carcinoma, basal cell carcinoma, benatoma, choriocarcinoma, carcinoma, pineatoma, medullary carcinoma, benatoma, choriocarcinoma, ceminoma, choriocarcinoma, medullary carcinoma, bronchogenic carcinoma, renal cell carcinoma, medullary carcinoma, bronchogenic carcinoma, propriatora, pineatoma, medullary carcinoma, choriocarcinoma, pineatoma, menuroblastoma, acustic neuroma, kidney carcinoma, propriatora, neuroblastoma, retinoblastoma, leukaemia, cu disease, breast tumours such as ductal and lobular adenocarcinoma, colon adenocarcinoma, such carcinoma, acute and choriocarcinoma, propriatora, carcinoma, colon adenocarcinoma, such carcinoma, colon adenocarcinoma, soft tissue sarcomas, colon adenocarcinoma, small cell lymphomas (nodular and colon carcinoma, propriatora denocarcinoma, small cell lung carcinoma, bladder cell carcinoma, propriatora denocarcinoma, small cell lung carcinoma, bladder cell carcinoma, propriatora denocarcinoma, ovarian adenocarcinoma, bladder cell carcinoma, propriatora denocarcinoma, small cell lung carcinoma, bladder cell carcinoma, propriatora denocarcinoma, small cell in an example cromposarcinoma, propriatora seguence is a complementarity determining carcinoma propriatora seguence is a complementarity determining colon should be should be should be solution of binders to the colon should be should be should be solution of binders to the colon should be should be should be solution of binders to the antigen. Mouse clone 13 was found to show specific binding to SPPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            variety of cancers: astrocytoma, fibrosarcoma, myxosarcoma, liposarcoma, oligodendroglioma, ependymoma, medulloblastoma, primitive neural ectodermal tumour (PNET), chondrosarcoma, osteogenic sarcoma, pancreatic ductal adenocarcinoma, small and large cell lung adenocarcinomas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to antigen-binding fragments, which are specific for stress protein-peptide complexes (SPPCs) associated with tumours and cancer associated SPPCs. The cancer-specific SPPC complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; immunostimulation; antibody; H11; tumour; cancer; murine; antigen-binding fragment; stress protein-peptide complex; SPPC; CDR; complementarity determining region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        are useful for initiating cancer-specific immunogenic responses against
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                                                                                                                                                                     The present invention describes vaccines for generating an anti-idiotype CC response to an antigen. Vaccine (A), comprises, apart from a carrier, a CC first immunoglobulin (I), comprising a variable region (V), or its CC fragment that includes V, which is identical to a second immunoglobulin (CC (II), or its corresponding fragment, which can bind specifically to an antigen (Ag), except for one or more amino acid (aa) substitutions in V. CC These substitutions involve replacement of Cys residues involved in CC disulfide bridge formation by residues that do not contain a thiol group. CC (A) are used to raise an anti-idiotype response (preferably both cellular CC and humoral), particularly for treating a wide range of tumours (or grecancerous conditions) and infectious diseases (viral, bacterial, CC fungal, parasitic), specifically syphilis, gonorrhea, acquired immune cc defictency syndrome, malaria, Shigella, Salmonella, hepatitis A or C, C fungal, parasitic), specifically syphilis, gonorrhea, acquired immune cc defictency syndrome, malaria, Shigella, Salmonella, hepatitive bacterial confections, or pneumococcal infections. Antibodies raised against the cc vaccine may be recovered and administered to a second subject. (I) may also be used for treating autoimmune diseases and allergy. Modifying (II) conformational constraints and results in significantly better anti-idiotype responses. The present sequence represents a Mab 31.1 CDR
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   Query Match
Best Local Similarity
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                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vaccines for generating an anti-idiotype response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 55; 107pp; English.
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10-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9925379-A1
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derived per
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98US-0081403P.
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                                                                                                                                                  es. The present sequence peptide from an example
   100.0%;
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   Score
Pred.
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Pred. No.
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30; DB 2; Length 7;
No. 1.8e+06;
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                                                                                                                                              represents a Mab
from the present
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disease;
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                                                                                                                                                                 second member (M2) of the pair. This portion is not present in the natural CDR and MI is a cancer antigen, infectious disease agent antigen, a cellular receptor for an infectious agent (provided the binding site is not NANP or NVDP) or a member of a receptor-ligand pair. mIg (also related molecules containing the same V region), optionally coupled to a chemotherapeutic agent, are used in therapeutic and/or prophylactic compositions, or vaccines, against cancers, infectious or autoimmune diseases (rheumatoid arthritis, ulcerative colitis or psoriasis) or allergies. They can also be used as immunoassay reagents for diagnosing these conditions mIg can be used in human or veterinary medicine. mIg have higher affinity or specificity than native antibodies. Manipulation of CDR ensures binding specificity and avoids the unpredictable
                                     immunization and screening procedures currently used. It is now possible to generate antibodies against antigens that are inaccessible, by virtue of immune tolerance or cryptic expression. mIg neutralize antigens directly or they induce an anti-idiotype immune response. This sequence represents a fragment of modified immunoglobulin that immunospecifically binds to a binding pair in which the first member is the human colon carcinoma-associated protein antigen
                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel modified immunoglobulin (mig), or i fragment, that binds immunospecifically to one member (M1) of a bind pair comprises a variable domain (V) having at least one CDR (complementarity determining region) that contains a portion of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complementarity determining region; infectious disease agent antigen; cellular receptor; infectious agent; chemotherapeutic agent; vaccine; cancer; infectious disease; autoimmune disease; rheumatoid arthritis; ulcerative colitis; psoriasis; allergy; immunoassay reagent; diagnosis; veterinary medicine; antibody; immunisation; antibody; immune tolerance; human colon cancer-associated protein antigen; cryptic expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 6; Page 75; 123pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified immunoglobulins including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-357555/30
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10-APR-1998;
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98US-0081403P.
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Sequence

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A new contraceptive vaccine is described which comprises an antibody CC which has at least one complementary determining region (CDR) containing CC a peptide fragment of a protein associated with reproductive function CC capable of eliciting an anti-idiotype response. The antibody is modified CC furtrachain disulfide bonds are substituted for amino acids lacking a CC sulfhydryl group and incapable of forming such bonds e.g Alanine. The CC vaccine formulation is suitable for both males and females. The method is CC convenient, effective, does not pose health risks and is cheap. In a CC particular example, a modified antibody derived from the monoclonal CC antibody MAD31.1 which recognises an antigen on human colon carcinomas CC had alanine residues substituted for the cysteine residues at positions C2 and 92 of the heavy chain variable region and at positions C3 and 88 CC of the light chain variable region. The modified variable regions were CC designated 2CAVECOL1 and 2CAVECOL1. MAD31.1, modified MAD31.1 and CC Shotinylated speptides derived from the CDR's of MAD31.1 were injected into mice (See MAY83638-Y83643) in order to generate antisera against the CCC CDR regions of MAD31.1 and the antibodies. This peptide was designated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel vaccine composition useful as contraceptive comprises first immunoglobulin capable of inducing an anti-idiotype response and identical to second immunoglobulin having complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine; contraceptive; antibody; monoclonal antibody; MAD; disulphide bond; complementary determining region; CDR; regroducti: immune response; anti-idiotype response; light chain; heavy chain; variable region; SP-10 epitope.
                                                                                                                                                                                                                                                                                                                                                                                               Example 6.6; Page 40; 77pp; English
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ilarity 100.0%;
Conservative
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Pred. No. 1.8e+06;
pred. Mismatches 0; Indels
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Sequence 7

RESULT 15 AAG80311 ID AAG80

AAG80311 standard; peptide; 10

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Best Local (
                                                                                                                            The present sequence represents a heavy chain hypervariable region of the murine anti-vascular endothelial growth factor (anti-VEGF) antibody. The sequence is used to construct the humanised anti-VEGF antibody of the invention. The humanised antibodies are used to inhibit VEGF-induced angiogenesis, particularly for treating or preventing tumours (of any type) and retinal disorders (e.g. age-related macular degeneration or diabetic retinopathy). They can also be used to treat other conditions that involve angiogenesis, e.g. rheumatoid arthritis, psoriasis,
                                                                                           Sequence 10 AA;
                                                                                                                                                                                                                                                                New humanised antibody with affinity for vascular endothelial growth factor - for treatment of tumours, retinal disease and other angiogenic states, also related nucleic acid, vectors and transformed cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Heavy chain hypervariable region; murine; humanised antibody; anti-vascular endothelial growth factor antibody; anti-VEGF a VEGF-induced angiogenesis; tumour; retinal disorder; age-related macular degeneration; diabetic retinopathy;
                                                                                                                    atherosclerosis,
                                                                                                                                                                                                                                                                                                                                       Baca M,
                                                                                                                                                                                                                                                                                                                                                                                       07-APR-1997;
06-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-VEGF antibody heavy chain hypervariable region CDRH1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW70611;
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                                             Similarity 5; Conserv
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NYCMN 10
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                                               Conservative
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                                                                                                                                                                                                                                            79;
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97US-00908469.
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                                                                                                                    Grave's disease, etc
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                                                                                                                                                                                                                                          100pp; English.
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                                                        100.0%;
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                                                        Score 30; DB Pred. No. 9.1;
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Pred. No. 1.8e+06;
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                                               Mismatches
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                                               Indels
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AAG80311;

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Search completed: April 4, 2005, 15:47:07 Job time: 11.984 secs
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Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel heavy and light-chain polypeptides of chimeric and humanized antibodies against human tumor necrosis factor alpha for low-immunogenicity treatment of TNF-related diseases such as toxic shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TNP-alpha; tumour necrosis factor-alpha; antibody; anti-human; CDR; heavy chain; light chain; complementarity determining region; vasotropic; antiarthritic; neuroprotective; protozoacide; toxic shock syndrome; rheumatoid arthritis; malaria; multiple sclerosis.
                                                                                                                                                                                                                 Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim la; Page 25; 36pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-066345/09.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-human TNF-alpha CDR-H1 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     syndrome.
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                                                                                                                                        Local Similarity hes 5; Conserv
                                                                                                       1 NYGMN 5
                                                                      NYGMN 10
                                                                                                                                            Conservative
                                                                                                                                          100.0%; Score 30; DB 5; Length 10; 100.0%; Pred. No. 9.1; tive 0; Mismatches 0; Indels
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Sequence 60, Appl Sequence 147, App Sequence 147, App Sequence 147, App Sequence 147, Appl Sequence 17, Appl Sequence 17, Appl Sequence 18, Appl

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No.
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Maximum Match 100%
Listing first 45 su
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                   Score
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seq length: 2000000000
        Published Applications AA:*

1: /cgn2=6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2=6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

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4: /cgn2=6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

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6: /cgn2=6/ptodata/1/pubpaa/US08 NEW PUB.pep:*

7: /cgn2=6/ptodata/1/pubpaa/US08 NEW PUB.pep:*

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12: /cgn2=6/ptodata/1/pubpaa/US09 NEW PUB.pep:*

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10: /cgn2=6/ptodata/1/pubpaa/US10D.PUB.pep:*

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  9 US-09-965-09-8

10 US-09-791-551-74

US-10-051-852-8

14 US-10-071-962-15

US-10-430-176-8

5 US-10-377-121-26

US-10-377-121-26

US-09-056-160B-1

4 US-10-234-671-1

6 US-10-018-245A-1

10S-10-723-434-111

5 US-10-723-434-111

5 US-10-723-434-112

1 US-10-723-434-116

1 US-10-723-434-116
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                                                                                                                                                                                                                                             SUMMARIES
Sequence 8, Appli
Sequence 74, Appli
Sequence 8, Appli
Sequence 15, Appli
Sequence 26, Appli
Sequence 17, Appli
Sequence 17, Appli
Sequence 1, Appli
Sequence 11, Appli
Sequence 111, Appli
Sequence 112, App
Sequence 112, App
Sequence 112, App
Sequence 112, App
                                                                                                                                                                                                  Description
                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1
US-09-965-099-8
; Sequence 8, Application US/09965099
; Patent No. US20020136725A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/965,099
FILING DATE: 26-Sep-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/346,487
FILING DATE: 40thrown>
PRIOR APPLICATION UNMBER: 09/346,487
FILING DATE: 40thrown>
ATTORNEY/AGENT INFORMATION:
NAME: Baumelster; Kirk
REGISTRATION UNMBER: 33,833
REFERENCE/DOCKET NUMBER: 50-438-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation STREET: 709 Swedeland Road CITY: King of Prussia
STATE: PA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ANTICOAGULANT AGENTS USEFUL TREATMENT OF THROMBOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Blackburn, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Feuerstein,
Patel, Arunb
                                                                                                                                                                                                                                                                                                                                                                                                  Arunbha:
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14 US-10-243-130-17
14 US-10-243-130-18
17 US-10-901-650-17
17 US-10-901-650-18
17 US-10-901-650-19
15 US-10-424-599-268610
18 US-09-965-099-84
18 US-10-051-852-84
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1 US-10-95-95-099-20

2 US-10-951-852-20

2 US-10-430-176-20

2 US-09-971-543-9

2 US-10-138-727A-2

3 US-10-138-727A-17

3 US-10-138-727A-18

4 US-10-138-727A-19

2 US-10-138-727A-19

2 US-10-138-727A-21

2 US-10-138-727A-21

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US-09-954-385-147
US-10-912-512-147
US-10-235-043-147
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Sequence 84, Appl Sequence 84, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 9, Appli Sequence 10, Appli Sequence 10, Appli Sequence 2, Appli Sequence 17, Appli Sequence 17, Appli Sequence 18, Appli Sequence 19, Appli Sequence 20, Appli Sequence 21, Appli Sequence 21, Appli Sequence 21, Appli Sequence 22, Appli Sequence 23, Appli

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RESULT 3
US-10-051-852-8
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                                                                                                                                                                                                      Sequence 8, Application US/10051852
Publication No. US20020146411A1
                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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SEQ ID NO 74
                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/791,551
CURRENT FILING DATE: 2001-02-26
PRIOR APPLICATION NUMBER: 60/185,390
PRIOR FILING DATE: 2000-02-28
PRIOR PPLICATION NUMBER: 60/233,625
PRIOR PPLICATION NUMBER: 60/233,625
PRIOR FILING DATE: 2000-09-18
NUMBER OF SEQ ID NOS: 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KLOETZER, WILLIAM S.
APPLICANT: HANNA, NABIL
TITLE OF INVENTION: METHOD FOR PREPARING ANTI-MIF ANTIBODIES
FILE REFERENCE: 037003/0277869
                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 5
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HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
NUMBER OF
                                  TITLE OF
                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                              Conservative
            INVENTION: ANTICOAGULANT AGENTS USEFUL IN TREATMENT OF THROMBOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                         Elackburn, Michael Church, William Gross, Mitchell Feuerstein, Giora Nichols, Andrew Padlan, Eduardo Patel, Arunbhai Sylvester, Daniel
SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 30; DB 10; 100.0%; Pred. No. 1.3e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 30; DB 9; Length 5; 100.0%; Pred. No. 1.3e+06;
                                                                                                                                                                                                                                                                                                                                                                          0,
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                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         DB 10; Length 5;
                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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CURRENT APPLICATION NUMBER: US/10/071,962
CURRENT FILING DATE: 2002-02-08
PRIOR APPLICATION NUMBER: US/09/303,155A
PRIOR PILING DATE: 1999-04-30
PRIOR APPLICATION NUMBER: 60/083,575
PRIOR FILING DATE: 1998-04-30
NUMBER OF SEQ ID NOS: 27
NUMBER OF SEQ ID NOS: 27
SEQ ID NO 15
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-071-962-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; SEQUENCE DESCRIPTION: SEQ ID NO: 8: US-10-051-852-8
                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15, Application US/10071962 Publication No. US20030170237A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 30; DB 13; Best Local Similarity 100.0%; Pred. No. 1.3e+06; Matches 5; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                  APPLICANT: Baufu Ni
APPLICANT: Bill N.C. Sun
APPLICANT: Cedily R.Y. Sun
TITLE OF INVENTION: G-CSF Receptor Agonist Antibodies and
TITLE OF INVENTION: Screening Method Therefor
FILE REFERENCE: 98-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/344,050
FILING DATE: 25-JUN-1999
APPLICATION NUMBER: 08/783,853
FILING DATE: 16-JAN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FRAGMENT TYPE: internal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 19406
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
CITY: King of Prussia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Baumeister, Kirk
REGISTRATION NUMBER: 33,833
REFERENCE/DOCKET NUMBER: P50438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/051,852
FILING DATE: 17-Jan-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ELEPHONE: 610-270-5096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 5;
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Query Match
Best Local Similarity
Thes 5; Conserve
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US-10-430-176-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: PRT
; ORGANISM: mouse
US-10-071-962-15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-377-121-26
Sequence 26, Application US/10377121
Sequence 26, Application US/103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-430-176-8
                      APPLICANT: GOVINDAM, SERENGULAM
APPLICANT: QU, ZHENGXING
APPLICANT: HANSEN, HANS
APPLICANT: GOLDENBERG, DAVID
TITLE OF INVENTION: RS7 ANTIBODIES
FILE REFERENCE: 018733/1163
CURRENT APPLICATION NUMBER: US/10/377,121
CURRENT FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: 60/360,299
PRIOR FILING DATE: 2002-03-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8, Application US/10430176
Publication No. US20030235587A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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CURRENT FILING DATE: 2003-05-05
PRIOR APPLICATION NUMBER: 09/817,960
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 09/359,202
PRIOR PILING DATE: 1999-07-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 111
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/010,018 PRIOR PILING DATE: 199-01-17 PRIOR APPLICATION NUMBER: 60/029,119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: 08/783,853
PRIOR FILING DATE: 1997-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: ANTICOAGULANT AGENTS USEFUL IN TREATMENT TITLE OF INVENTION: OF THROMBOSIS FILE REFERENCE: P50816-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 10/051,852
PRIOR FILING DATE: 2002-01-17
PRIOR APPLICATION NUMBER: 09/344,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/095,714 PRIOR FILING DATE: 1998-08-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1996-10-24
NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Feuerstein, Giora 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Home sapins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No. 1.3e+06;
0; Mismatches 0;
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                                                                                                                         US-09-056-160B-1
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                                                         Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09056160B Patent No. US20020032315A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION UNMBER: US/09/056,160B

FILING DATE: 06-Apr-1998

CLASSIFICATION: 424

CDASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                  TELECOMMUNICATION INFORMATION: 650/225-1896
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                            FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E
REGISTRATION NUMBER: 28,61
REFERENCE/DOCKET NUMBER: F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 5
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1 DNA Way
CITY: South San F:
STATE: California
                                                           Local Similarity
mes 5; Conserv
                                                                                                                                                      LENGTH: 10 amino acids TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
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 6 NYGMN 10
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                        NYGMN 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wells, James A.
Presta, Leonard G
Lowman, Henry B.
Chen, Yvonne M.
                                                         100.0%; Score 30; DB 9; ilarity 100.0%; Pred. No. 6.2; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Baca, Manuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                                                                                                                                                                28,616
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 30; DB 15;
Pred. No. 1.3e+06;
Mismatches 0;
                                                                                        Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 5;
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                                                            Сарв
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RESULT 8 US-10-234-671-1 ; Sequence 1, Application US/10234671

Publication No. US20030190317A1 GENERAL INFORMATION:

APPLICANT: Baca, Manuel

Wells, James A. Presta, Leonard (Lowman, Henry B.

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Yvonne M.

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                                                                                                                                                                                                                                                  Sequence 1, Application US/10018245A Publication No. US20040115196A1 GENERAL INFORMATION:
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Best Local
             CURRENT APPLICATION NUMBER: US/10/018,245A CURRENT FILING DATE: 2002-07-12 PRIOR APPLICATION NUMBER: UF 117394/2000 PRIOR FILING DATE: 2000-04-19 NUMBER OF SEQ ID NOS: 19
                                                                                                                      APPLICANT: FUKUDA, Yoshiaki
APPLICANT: NAGAHIRA, Kazuhiro
APPLICANT: NAKANISHI, Toshihiro
APPLICANT: NAKANISHI, Toshihiro
TITLE OF INVENTION: Novel recombinant antibody, amino acid sequences of its complemen
TITLE OF INVENTION: determining regions and genes encoding the same
FILE REFERENCE: 46224
SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 1:
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TELEPHONE: 650/225-8674
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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FILING DATE: 06-APR-1998
APPLICATION NUMBER: 60/126446
FILING DATE: 07-APR-1997
APPLICATION NUMBER: 60/054856
FILING DATE: 06-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                          NYGMN 10
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REFERENCE/DOCKET NUMBER: P1093R2C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/234,671 FILING DATE: 03-Sep-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
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Pred. No. 6.2;
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RESULT 11
US-10-723-434-112
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Best Local Similarity
Thehes 5; Conserve
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US-10-018-245A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/284,407
PRIOR FILLING DATE: 2001-04-17
PRIOR PELLCATION NUMBER: US 10/125,687
PRIOR FILLING DATE: 2002-04-17
PRIOR FILLING DATE: 2002-04-17
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR FILLING DATE: 2002-05-20
PRIOR PILLING DATE: 2003-05-20
PRIOR FILLING DATE: 2003-05-20
NUMBER OF SEQ ID NOS: 156
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                                                                                                                                                            GENERAL INFORMATION:
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Best Local
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APPLICANT: Li, Yan
TITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST VASCULAR ENDOTHELIAL GROWTH FACTOR
FILE REFERENCE: 26050-709.501
CURRENT APPLICATION NUMBER: US/10/723,434
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 60/284,407
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                                                                                                            APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/723,434
CURRENT FILING DATE: 2003-11-26
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TYPE: PRT
ORGANISM: mouse
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TITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST VASCULAR ENDOTHELIAL GROWTH FACTOR
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                            Wang, Kevin C.
Hsieh, Mark
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Hsieh, Mark
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o. US20040133357A1
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tive 0;
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Pred. No.
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Pred. No. 6.2;
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APPLICANT: Luo, Peizhi
APPLICANT: Luo, Peizhi
APPLICANT: Wang, Kevin C.
APPLICANT: Wang, Kevin C.
APPLICANT: Wang, Kevin C.
APPLICANT: Wang, Kevin C.
APPLICANT: Haich, Mark
APPLICANT: Li, Yan
TITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST VASCULAR ENDOTHELIAL GROWTH FACTOR
FILE REFERENCE: 2605-709.501
CURRENT APPLICATION NUMBER: US/10/723,434
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 60/284,407
PRIOR APPLICATION NUMBER: US 10/125,687
PRIOR APPLICATION NUMBER: US 10/125,687
PRIOR FILING DATE: 2002-04-17
PRIOR FILING DATE: 2002-05-20
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-05-20
NUMBER OF SEQ ID NOS: 156
RESULT 13
US-10-723-434-122
; Sequence 122, Application US/10723434
; Publication No. US20040133357A1
; GENERAL INFORMATION:
; APPLICANT: Zhong, Pingyu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2001-04-17
PRIOR APPLICATION NUMBER: US 10/125,687
PRIOR PILING DATE: 2002-04-17
PRIOR PILING DATE: 2002-05-20
PRIOR PILING DATE: 2002-05-20
PRIOR APPLICATION NUMBER: US 10/443,134
PRIOR PILING DATE: 2003-05-20
NUMBER OF SEQ ID NOS: 156
SOFTWARE: PatentIn version 3.1
SEQ ID NO 112
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Best Local Similarity
Marches 5; Conserve
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US-10-723-434-116
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SEQ ID NO 116
LENGTH: 10
TYPE: PRT
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                                                                                                                                                                                                                             Matches
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OTHER INFORMATION: VH/CDR1
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TYPE: PRT
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o. US20040133357A1
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Pred. No. 6.2;
0; Mismatches
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Pred. No.
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6.2;
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; OTHER INFORMATION: 725.1 VH1 CDR US-10-687-035-60
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Best Local Similarity
Thes 5; Conserve
RESULT 15
US-09-954-385-147
; Sequence 147, Application US/09954385
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CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 60/284,407
PRIOR FILING DATE: 2001-04-17
PRIOR APPLICATION NUMBER: US 10/125,687
PRIOR FILING DATE: 2002-04-17
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR FILING DATE: 2002-05-20
PRIOR APPLICATION NUMBER: US 10/443,134
PRIOR FILING DATE: 2003-05-20
PRIOR PRIOR DATE: 2003-05-20
PRIOR PRIOR DATE: 2003-05-20
NUMBER OF SEQ ID NOS: 156
                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/418,828
PRIOR FILING DATE: 2003-10-12
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 60
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SEQ ID NO 122
LENGTH: 10
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Best Local
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                                                                                                                                                                              Matches
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TITLE OF INVENTION: ANTIBODIES THAT BIND CELL-ASSOCIATED
TITLE OF INVENTION: CA 125/0772P AND METHODS OF USE THER
FILE REFERENCE: 6750-214-999
CURRENT APPLICATION NUMBER: US/10/687,035
CURRENT FILING DATE: 2003-10-15
PRIOR APPLICATION NUMBER: 60/485,986
PRIOR FILING DATE: 2003-07-10
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TITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST VASCULAR ENDOTHELIAL GROWTH FACTOR
FILE REFERENCE: 26050-709.501
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                                                                                                                                                                                                                                                                                        LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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OTHER INFORMATION: VH/CDR1
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Hsieh, Mark
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GENERAL INFORMATION:
APPLICANT: Holdwin, Toby L.
APPLICANT: Wan Gastel, Franciscus J.C.
APPLICANT: Wan Gastel, Franciscus J.C.
APPLICANT: Warnes, Christopher J.
APPLICANT: Warney, Christopher J.
APPLICANT: Warne
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A;Cross-references: GB:X03301; NID:g51757; PIDN:CAA27040.1; PID:g773215 A;Note: this sequence was determined from the differentiated gene C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
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C;Species: Mus musculus (house mouse)
C;Date: 19-Nov-1988 #sequence_revision 19-Nov-1988 #text_change 23-Jul-1999
C;Accession: D24672
R;Winter, E.; Radbruch, A.; Krawinkel, U.
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Ig heavy chain V region (subgroup XI) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 20-Feb-1995 #sequence revision 21-Nov-1998 #text_C;Accession: S24765; S24773; S24778 R;Klages, S. submitted to the EMBL Data Library, August 1992 A;Reference number: S24763
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Reference number: S24776
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R;Thomas, J.W.
submitted to the EMBL Data Library, August 1992
A;Reference number: S24776
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C;Accession: S26325
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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S19968
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A; Residues: 1-105 < THO>
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A; Residues: 1-105 < KLW >
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A;Molecule type: DNA
A;Residues: 1-105 <KLA>
A;Cross-references: EMBL:Z15001
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A;Molecule type: mRNA
A;Residues: 1-115 <WEI's
A;Cross-references: UNIPROT:Q921A6; EMBL:X65089
C;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                 A; Description: Structural A; Reference number: S19963 A; Accession: S19968
                                                                                                                                                            R; Weissenhorn, W.; Riethmueller, G.; Weiss, E.; Submitted to the EMBL Data Library, March 1952
                                                                                                                                                                                                        Ig heavy chain V region (M-T408) - mouse (fragment)
C;Species: Mus muscultus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 09-Jul-2004
C;Accession: S19968
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                                                                              A; Status: preliminary
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les 5; Conservative
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                                                                                                                                             Structural characterization
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                                                                                                                                                                                         Riethmueller, G.; Weiss, E.M.; Rieber,
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Pred. No. 12;
0; Mismatches
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                                                                                                                                               of CD4 mAb
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RESULT 10
819967
Ig heavy c
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C;Bspecies: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change
C;Accession: $19965
C;Accession: $19965
C;Accession: $19965
C;Accession: $19963
C;Accession: $19963
C;Accession: $19963
C;Accession: $19963
A;Reference number: $19963
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A;Residues: 1-116 <PAU>
A;Cross-roferences: UNIPROT:O13541;
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                   N;Alternate names: hypothetical protein L8003.10-b
C;Species: Saccharomyces cerevisiae
C;Date: 20-Jul-1996 #sequence_revision 23-Aug-1996 #text_change 09-Jul-2004
C;Accession: S69305
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C;Superfamily: immunoglobulin V region; immunog.
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immonoglobulin
F;8-91/Domain: immunoglobulin homology <IMM>
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A;Status: preliminary
A;Molecule type: mRNA
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F;10-93/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                              Gene: MIPS:YLR280C
;Cross-references: SGD:S0004270
;Map position: 12R
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Best Local
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5; Conserv
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llarity 100.0%;
Conservative 0
                                                                                                                                                      Conservative
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   (M-T406) - mouse (fragment)
                                                                                                                                                                  100.0%;
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100.0%; Pred. No. 13;
tive 0; Mismatches
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RESULT 11
A53285
Ig heavy chain V and J regions, monoclonal antibody SCET.M8.1 - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 20-Jun-2000
C;Accession: A53285
R;Sawada, J; Mizusawa, S; Terao, T.; Naito, M.; Kurosawa, Y.
Mol. Immunol. 28, 1063-1072, 1991
A;Title: Molecular characterization of monoclonal anti-steroid antibodies: prima and their pH-reactivity profiles.
A;Reference number: A53285; MUID:92017897; PMID:1922102
A;Accession: A53285
A; ACCEBBANCE PREJIMINATY
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-120 <FEL>
A; Residues: 1-120 <FEL>
A; Cross-references: GB: M90690; NID: g195065; PIDN: AAA38146.1; PID: g195066
A; Cross-references: GB: M90690; NID: g195065; PIDN: AAA38146.1; PID: g195066
A; Cross-reference extracted from NCBI backbone (NCBIN: 109960, NCBIP: 109961)
                                                                                                                                                                                     L6 mAb heavy chain V region - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Date: 27-Apr-1993 #sequence revision 18-Nov-1994 #text_change 21-Jan-2000
C; Accession: 842848; 533903
R; Fell, H.P.; Gayle, M.A.; Yelton, D.; Lipsich, L.; Schieven, G.L.; Marken, J. Biol. Chem. 267, 15552-15558, 1992
A; Title: Chimeric L6 anti-tumor antibody. Genomic construction, expression, A; Reference number: A42848; MUID:92348410; PMID:1639794
A; A; A; Ccession: 842848
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C;Date: 06-Peb-1995 #sequence_revision 06-Peb-1995 #text_change 09-Jul-2004
C;Accession: S19967
R;Weissenhorn, W., Riethmueller, G.; Weiss, E.M.; Rieber, E.P.
submitted to the EMBL Data Library, March 1992
A;Description: Structural characterization of CD4 mAb.
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C;Keywords: heterotetramer; immunoglobulin
F;12-95/Domain: immunoglobulin homology <IMM>
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A;Molecule type: mRNA
A;Residues: 1-118 <WEI>
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F;15-98/Domain: immunoglobulin homology <IMM>
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A; Accession: S19967
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ilarity 100.0%;
Conservative 0
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larity 100.0%;
Conservative 0,
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Pred. No. 13;
0; Mismatches
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Pred. No. 13;
0; Mismatches
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C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin_____
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig heavy chain V region (clone micro m+ 46-12) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 21-Jan-2000
C;Accession: PH1004; PH1406
R;Shirasawa, T.; Miyazoe, I.; Hagiwara, S.; Kimoto, H.; Shigemoto, K.; Taniguchi, M.;
J. Exp. Med. 176, 1209-1214, 1992
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PH1404
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A, Molecule type: mcNA
A, Molecule type: mcNA
A;Residues: 1-120 <WER>
A;Cross-references: GB:X55179; NID:g56471; PIDN:CAA38964.1; PID:g56472
A;Cross-reference was determined from the germline gene
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
A; Residues: 115-121 <SH2>
                                                                                                                                                                                                                                                                                                                                                      A;Reference number: PH1403; MUID:93018837; PMID:1402663
A;Accession: PH1404
A;Molecule type: DNA
A;Residues: 1-124 <SHI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Title: Heavy chain variable (VH) region diversity generated by VH gene replacement in
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A;Residues: I-120 <FE2>
A;Cross-references: EMBL:M90691
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;15-98/Domain: immunoglobulin homology <IMM>
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Ig heavy chain precursor V region - synthetic C;Species: synthetic C;Species: synthetic C;Date: 13-Jan-1995 #sequence_revision 30-Apr-1998 #text_change 20-Oct-2000 C;Accession: S33905 R;Liu, A.Y.; Robinson, R.R.; Hellstroem, K.E.; Murray Jr., E.D.; Chang, C.P.; Hellstroem, Proc. Natl. Acad. Sci. U.S.A. 84, 3439-3443, 1987 Proc. Natl. Acad. Sci. U.S.A. 84, 3439-3443, 1987 A;Title: Chimeric mouse-human IgG1 antibody that can mediate lysis of cancer cells. A;Reference number: S33905; MUID:87204152; PMID:3106970 A;Accession: S33905
Search completed: April Job time: 2.80511 secs
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A;Residues: 1-146 <LIU>
A;Cross-references: EMBL:M16072; NID:g195270; PIDN:AAA38229.1; PID:g195271
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		Query Matc Best Local Matches	SEQUENCE Cherry J. Submitted EMBL; U1: PIR; S693 SEQUENCE	SEQUENCE Pauley A. Submitted [3] SEQUENCE Waterston Submitted [4]	Entian Entian Heuss- Louis Muller Muller Portet Scharf Underw Vieren Wedler	Name=YLR2 Name=YLR2 Saccharom Eukaryota Saccharom NCBI_TaxI [1] SEQUENCE MEDLINE=9 Johnston	ESULT 1 13541 D 013541 C 013541; C 01-JAN-1 T 01-JAN-1 T 01-MAR-2 T 11-MAR-2		S S S S S S S S S S S S S S S S S S S
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OrderedLocusNames=BCE0627;
Bacillus cereus (strain ATCC 10987).
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.
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Imidazole glycerol phosphate synthase subunit hisH (EC 2.4.2.-) (IGP synthase glutamine amidotransferase subunit) (IGP synthase subunit hisH) (ImGP synthase subunit hisH)
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PROSITE; PS00442; GATASE_TYPE_T; 1.
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Bacteria, Proteobacteria, Gammaproteobacteria, Enterobacteriales,
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                                                                                                                           "Complete genome sequence of the industrial bacterium Bacillus licheniformis and comparisons with closely related Bacillus species."; Genome Biol. 5:R77-R77(2004).
EMBL; AE01733; AAU40213.1; -.
EMBL; CP000002; AAU22864.1; -.
                                                                                                                                                                                        sorokin A., Bolotin
Berka R.M.;
                                                                                                                                                                                                                                                                                               Veith B., Herzberg C., Steckel S., Feesche J., Maurer K.H., Ehrenzeich P., Baeumer S., Henne A., Liesegang H., Merkl R., Ehrenzeich A., Gottschalk, G., Ehrenzeich A., Gottschalk Grene Sequence of Bacillus licheniformis DSM13, "The Complete Genome Sequence of Bacillus licheniformis DSM13, Organism with Great Industrial Potential.";
J. Mol. Microbiol. Biotechnol. 7:204-211(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PubMed=14960714; DOI=10.1093/nar/gkh258;
Rasko D.A., Ravel J., Oekstad O.A., Helgason E., Cer R.Z., Jiang L., Shores K.A., Fouts D.E., Tourasse N.J., Angiuoli S.V., Kolonay J.F., Nelson W.C., Kolstoe A.-B., Fraser C.M., Read T.D.;
"The genome sequence of Bacillus cercus ATCC 10987 reveals metabolic adaptations and a large plasmid related to Bacillus anthracis pXO1."; Nucleic Acids Res. 32:977-988(2004).
                                                                                                     Hypothetical SEQUENCE 24
                                                                                                                                                                                                                Rey M.W., Ramaiya P., Nelson B.A., Brody-Karpin S.D., Zaretsky Tang M., de Leon A.L., Xiang H., Gusti V., Clausen I.G., Olsen Rasmussen M.D., Andersen J.T., Jorgensen P.L., Larsen T.S.,
                                                                                                                                                                                                                                                                                                                                                                          STRAIN=DSM 13;
PubMed=15383718;
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Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus
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"Molecular phylogeny and divergence time estimates
groups: evidence from multiple genes.";
Mol. Biol. Evol. 18.777-791(2001).

EMBL; AP332021; AAK71897.1;
GO; GO:0004872; F:receptor activity; IEA.
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Steppan S.J., Adkins R.M., Anderson J.;

Submitted (MAY-2003) to the EMBL/GenBank/DDBJ

EMBL; AY29493; AAQ62681.1; -.

GO; GO:0004872; F:receptor activity; IEA.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Scivrognathi; Muridae; Gerbillinae;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Acomys.
NCBI_TaxID-60745;
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llarity 100.0%;
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Pred. No. 1.5
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Pred. No. 1.7e+02;
Mismatches 0;
                                   PRT;
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Q924G5
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05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
Growth hormone receptor ()
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NON TER
SEQUENCE
Q924G5 PRELIMINARY; PRT; 302 AA. Q924G5.
Q1-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Growth hormone receptor (Fragment).
Name=GHR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Receptor.
NON_TER
NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Steppan S.J., Adkins R.M., Anderson J.; Submitted (MAY-2003) to the EMBL/GenBank/EMBL; AY294905; AAQ62663.1; GO; GO:0004872; F:receptor activity; IBA.
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05-UUL-2004 (TrEMBLrel. 27, Last sequence up
05-UUL-2004 (TrEMBLrel. 27, Last annotation
Growth hormone receptor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10085;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cricetomys gambianus (Gambia rat).
Eukaryota; Metazoa; Chordata; Cran
Mammalia; Eutheria; Rodentia; Sciu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Name=GHR;
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Beamys hindei (long-tailed pouched rat).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetomyinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                            200 NYGMN 204
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301 AA;
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27, Last sequence update)
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: (Fragment).
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e EMBL/GenBank/DDBJ databases.
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Pred. No. 1.7e+02;
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Pred. No. 1.7e+02;
; Mismatches 0;
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; Cricetomyinae;
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RESULT 10
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Q9CLC4
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Q9CLC4;
01-JUN-2001
01-JUN-2001
01-JUN-2003
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Q895E8;
01-JUN-2003
01-JUN-2003
01-JUN-2003
Name=corA;
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SEQUENCE
                     CorA.
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Hypothetical protein CTC01328.
OrderedLocusNames=CTC01328;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Adkins R.M., Gelke E.L., Rowe D., Honeycutt R.L.; "Molecular phylogeny and divergence time estimates groups: evidence from multiple genes."; Mol. Biol. Evol. 18:777-791(2001).

EMBL; AF332022; AAK71898.1; -.
EGG; GG.0004872; F:receptor activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
Gerbillurus.
                                                                                                                                                                                                                                                                                                                                                                                              Complete proteome.
SEQUENCE 305 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                tetanus disease.";
Proc. Natl. Acad. Sci. U.S.A. 100:1316-1321(2003).
EMBL; AE015940; AA035892.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Decker I., Her Gottschalk G.;
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Decker I., Herzh
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MEDLINE=22457253; PubMed=12552129;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clostridium tetani.
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[1]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genome sequence of Clostridium tetani,
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                                                                                                                                                                                                                             NYGMN
OrderedLocusNames=PM1315;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    457253; PubMed=12552129; DOI=10.1073/pnas.0335853100;
n H., Baeumer S., Fricke W.F., Wiezer A., Liesegang F
Herzberg C., Martinez-Arias R., Merkl R., Henne A.,
                                  (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                                                                                         PRELIMINARY;
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Rodentia;
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SEQUENCE
                                                                                                                                                                                                                                           MEDIZINE-20188199; PubMed=11823852; DOI=10.1038/415497a; MEDIZINE-21681879; PubMed=11823852; DOI=10.1038/415497a; Salanoubat M., Genin S., Artiguenave F., Gouzy J., Mangenot S Arlat M., Billault A., Brottier P., Camus J.C., Cattolico L., Chandler M., Choisen N., Claudel-Renard C., Cunnac S., Demang, Gaspin C., Lavie M., Moisan A., Robert C., Saurin W., Schiex Siquier P., Thebault P., Whalen M., Wincker P., Levy M.,
                                                                                                                                        EMBL; AL646080; CAB17816.1; ...
GO; GO:0016020; C:membrane; IEA.
GO; GO:0046873; F:metal ion transporter activity;
GO; GO:0030001; P:metal ion transport; IEA.
                                                                                                                                                                                                  Gaspin C., Lavie M., Moisan A., Robert C., Saurin W., Schiex T., Siguiler P., Thebault P., Whalen M., Wincker P., Levy M., Weissenbach J., Boucher C.A.;
"Genome sequence of the plant pathogen Ralstonia solanacearum.";
Nature 415:497-502(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-UIN-2003 (TrEMBLrel. 24, Last annotation update)
PROBABLE MAGNESIUM AND COBALT TRANSPORT TRANSMEMBRANE
Name=RS01784; OrderedLocusNames=RSp0665;
                                                                                                                               Pfam; PF01544; CorA;
                                                                                                                                                                                                                                                                                                                                      STRAIN=GMI1000;
                                                                                                                                                                                                                                                                                                                                                                                                                       Plasmid megaplasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                      Ralstonia solanacearum (Pseudomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GO; GO:0015693; P:magnesium ion transport; InterPro; IRR004488; CorA transp. InterPro; IPR005253; Mg2+_transptCorA. Pfam; PF01544; CorA; 1. TIGRPAMS; TIGR00383; corA; 1.
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EMBL; AE006170; AAK03399.1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0015087; F:cobalt ion transporter activity; IEA.
GO; GO:0015095; F:magnesium ion transporter activity; IEA.
GO; GO:0006824; P:cobalt ion transport; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                            Burkholderiaceae;
                                                                                                                                                                                                                                                                                                                                                                                                            Bacteria;
                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=305;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          May B.J., Zhang Q., Li L.L., Paustian M.L., Whittam "Complete genomic sequence of Pasteurella multocida Proc. Natl. Acad. Sci. U.S.A. 98:3460-3465(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=21145866; PubMed=11248100; DOI=10.1073/pnas.051634598;
May B.J., Zhang Q., Li L.L., Paustian M.L., Whittam T.S., Kapu
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323 AA;
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                                            Conservative
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100.0%; F
tive 0;
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                                          Score 30; DB
Pred. No. 1.8
0; Mismatches
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Pred. No. 1.8e+02;
Mismatches 0;
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                                                                    Length 323;
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RESULT 13
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Q9E3T9
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Q9E3T9;
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085033; PRELIMINARY;
01-NOV-1996 (TrEMBLrel. 01;
01-NOV-1996 (TrEMBLrel. 20;
01-MAR-2002 (TrEMBLrel. 20;
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GO; GO:0019028; C:viral capsid; IEA.
InterPro; IPRO1963; VP7.
Pfam; PF00434; VP7; 1.
ProDom; PD000191; VP7; 1.
SEQUENCE 326 AA; 37318 MW; ED237
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01-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                       Ciarlet M., Ludert J.E., Liprandi F.;
Submitted (AUG-1994) to the EMBL/GenBank/DDBJ databases.
EMBL; L35057; AAA47091.1; -.
GO; GO:0019028; C:viral capsid; IEA.
InterPro; IPR001963; VP7.
Pfam; PF00434; VP7; 1.
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                                                                                                                                                                                                                                                                                                                                      Prodom; PD000191; VP7; 1.
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Viruses; dsRNA viruses; Reoviridae;
NCBI_TaxID=10913;
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Viruses; dsRNA viruses; Reoviridae; Rotavirus; Human rotavirus
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Local Similarity 100.0%;
les 5; Conservative 0
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Pred. No. 1.8e+02;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence upda
01-MAR-2003 (TrEMBLrel. 23, Last annotation up
Outer capsid protein VP7.
Human rotavirus A.
Viruses; dsRNA viruses; Reoviridae; Rotavirus.
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193.690 Million cell updates/sec
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ALIGNMENTS	AAB84740	ADC27449	ADC27441	ADH62638	ADH62639	AAY83642	AAY08699	AAY17490	ADS87337	AAG80312	AAW85692	AAW85691	AAW85689	AAW85688	AAW85693	AAW85694	AAW85695	AAB36825	AAB12562	ADC65008
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AAY90823 standard; protein; 115 B

AAY90823;

29-AUG-2000 (first entry)

520C9 hybridoma VH domain SEQ ID NO:24.

Antigen binding site; immunoglobulin; cancer antigen; immunological; antibody; tumour; human; mucin; cancer; cytostatic; hybridoma; specific binding assay; affinity purification; drug targeting; toxin targeting; imaging; genetic; therapeutic.

Homo sapiens

US6054561-A.

25-APR-2000

08-FEB-1984; 07-JUN-1995; 84US-00577976 95US-00483749.

11-JAN-1985; 21-MAR-1986; 08-MAY-1988; 11-AUG-1994; 86US-00842476. 88US-00190778. 94US-00288981.

(CHIR) CHIRON CORP.

Ring

WPI; 2000-338508/29. N-PSDB; AAA38907.

treating Monoclonal antibody capable of binding useful for affinity purification, drug cancer. 유당 human breast cancer antigen targeting, imaging, and

Disclosure; Fig 12; 57pp; English.

RRESULT 1
RAYSORS 1
RAYSOR The present invention describes a monoclonal antibody (MAb) (I) that binds to a human breast cancer antigen that is also bound by MAb 454C11 and 520C9 (produced hybridoma ATCC HB8484 and HB8696, respectively). Also described is a hybridoma that produces (I). (I) is useful in specific binding assays, affinity purification, drug or toxin targeting, imaging, and genetic or immunological therapeutics for various cancers. The Also

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RESULT 2
AAW02280
ID AAW0
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Best Local S
Matches 17
           Variable heavy (VH) and variable light (VL) genes were cloned from a $2009 hybridoma cDNA library, using probes directed toward the antibody constant and joining regions. A two single chain Fr (SFV) gene was constructed by connecting the VH and VL genes with a Ser rich polypeptide linker. The resulting 52009 two sFV gene, which encodes the present sequence, was inserted into an expression vector, transformed into E coli, and protein expression induced by the addn. of IPTG to the culture medium. A compsn. comprising a carrier and the 2 sFV protein prod. can be used for in vivo imaging, and drug targetting experiments. The 2 sFV protein prod. is a homodimer, in which both fragments target the same antigen, therefore giving greater binding avoidity and longer tissue retention times, compared to individual special prod. fragments.
                                                                                                                                                                                                                                                                                                                                                              Compsns. contg. antigen-targetting antibody fragment constructs - comprising dimer of single-chain Fv fragments.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       520C9 anti-c-erbB-2 two single chain Fv construct.
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29-OCT-1996
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(Updated on 25-MAR-2003 to correct PF
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Sequence 243

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                                                                                                                                                                                                                                                                                     This represents the protein sequence of a 520C9 sFv' (single chain Fv) construct. This was constructed by connecting the Vh and Vl genes with a DNA sequence encoding a Serine rich linker. 520C9 is a monoclonal antibody useful in targeting c-erbB-2 antigen. This dimeric construct can be used in the methods of invention of imaging a preselected antigen expressed in a mammal. The methods are used in magnetic resonance imaging of c-erbB-2 or related antigens in cancer diagnosis. The biosynthetic constructs have enhanced properties as in vivo targetting agents in comparison with intact monoclonal antibodies or their Fab fragments. The dimeric constructs permit the in vivo targetting of an epitope on an antigen with greater apparent avidity, including greater tumour specificity, tumour localisation and tumour retention properties than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antigen tumour;
                                                                                                                                                                                                               Sequence 243 AA;
                                                                                                                                                                                                                                                                    that of the Fab fragment having the same CDRS as the construct
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07-OCT-1993;
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(CREA-) CREATIVE BIOMOLECULES
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17; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             imaging; single chain Fv; sFv; linker; dimeric; cancer; c-erbB-2;
diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV21798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                          WINTYTGOSTYADDEKE
WINTYTGQSTYADDFKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WINTYTGQSTYADDFKE
                                                                                                  100.0%; ilarity 100.0%; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ring DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-00831967.
93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95US-00461838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Huston JS,
66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  66
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                                                                                                                                Score 97;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 97; DB 2;
Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8
                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Houston LL;
                                                                                                                                1.1e-06;
                                                                                                                                                               DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                          Length 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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DT 15-J
CX CB
DE 520C
XX CC
DB 520C
XX CC
DB 67-V
XX CO
XX Unid
XX UN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents an antibody 520C9 sFV. Variable heavy and light sequences of antibody 7520C9 are connected, together with a serine linker, to produce the present single chain FV gene. The present sequence exemplifies the invention. Dimers of the single chain FV are used for targeted delivery of drugs or imaging agents (e.g. cytotoxins, prodrugs or 99m-technetium) to antigen-expressing cells, particularly for treatment or diagnosis of tumours (especially of ovary or breast)
                                                                                                                                                                                                                                                                            520C9 aFv protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABW00716 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid encoding single-chain Fv fragment specific for antigens and having C-terminal tail for crosslinking to form dimer with improved pharmacokinetic properties, used to deliver drugs and imaging agents,
    US2002168375-A1
                                                                                  Unidentified
                                                                                                                                                                 Cell proliferation; cytotoxic agent; drug targetting; sinçle-chain Fv;
sFv.
                                                                                                                                                                                                                                                                                                                                                               15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                             ABW00716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 243 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Col 35-36; 29pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          especially to tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-023541/02.
N-PSDB; AAV63399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-FEB-1992;
07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antigen-expressing cell.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US5837846-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             520C9 sFv; antigen; tumour cell; antibody 520C9; targeted delivery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           520C9 sFv sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW80424 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CREA-) CREATIVE BIOMOLECULES INC (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW80424;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ħ,
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                      (first entry)
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93US-00133804.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   243
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Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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Best Local S
Matches 17
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07-OCT-1993;
05-JUN-1995;
26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a binding protein formulation for targetting an epitope on an antigen expressed in mammal. The formulation comprises of dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain. The invention is useful for imaging a preselected antigen in a mammal expressing the antigen. It is also useful for targetting drugs that inhibits cell proliferation and cytotoxic agents that kill cells. The present sequence is 52009 sFv protein. This sequence is used in the exemplification of the
                          Huston JS,
                                                                                                                                                     06-FEB-1992;
                                                                                                                                                                                                       05-FEB-1993;
                                                                                                                                                                                                                                                          19-AUG-1993.
                                                                                                                                                                                                                                                                                                             WO9316185-A2
                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                          Single chain
biosynthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence of 520C9 sFv protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
07-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR39569;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR39569 standard; protein; 246 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 243 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 19-20; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel binding protein formulation for targeting epitope on antigen in mammal, comprising dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-765156/72.
N-PSDB; AAD61485.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUN-2001; 2001US-00887853.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Huston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ch 100.0%;
l Similarity 100.0%;
17; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       \vdash
                                                                          CREATIVE BIOMOLECULES CETUS ONCOLOGY CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WINTYTGOSTYADDFKE 17
                          Houston LL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Houston
                                                                                                                                                                                                                                                                                                                                                                                                          Fv polypeptide; VH-VL heterodimer; immunoglobulin; single polypeptide chain binding site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-00558741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                     9208-00831967
                                                                                                                                                                                                       93WO-US001055
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95US-00462641.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-00831967
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                     Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    66
                          BB,
                                                                                                  INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 97; DB 7; Pred. No. 1.1e-06; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽B,
                          Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the
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RESULT 7
AAU04944
ID AAU04944
ID AAU04944
AC AAU0
XX AAU0
DT 06-2
DT 24-C
XX Huma
XX Huma
XX Homc
OS Mus
Synt
COS Mus
Synt
FFT Regi
FT Regi
FT Pept
FT Regi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). A single chain Fv (EFV) is a covalently linked VH-VL heterodimer which is expressed from a gene fusion including VH- and VL- encoding genes connected by a peptide-encoding linker. Such linker sequences are set forth in AA residues 116-135 in AAR39569, which includes part of the 16 AA linker sequences in AAR39572. Using AAQ46084 for the 520C9 monoclomal antibody, a single chain polypeptide can be produced having a binding affinity for a c-erbB-2 related antigen. 'X' in AAR39569 refers to the location of a stop codon in AAQ46084. (Updated on 25-MAR-2003 to correct PN field.)
                                                       Region
                                                                                                                             Region
                                                                                                                                                                                                       Region
                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic. Chimeric.
                                                                                                                                                                                                                                                                           Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Humanised;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAU04944;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           c-erbB-2 refers to a protein antigen expressed on the surface cells, such as breast and ovarian tumour cells, which is an approximately contains the contains and contains the contains tha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Humanised anti-p185 single chain antibody, 520C9H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Page 60-61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New single chain Fv polypeptide binding to C-erbB-2 tumour antigen imaging or treating breast or ovarian cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAU04944 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB; AAQ46084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1993-272889/34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 WINTYTGOSTYADDFKE 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WINTYTGQSTYADDFKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibody; p185; 520C9H; interleukin-2; IL-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adenocarcinoma.
                                                         /label= CDR
/note= "Complementarity determining region"
'note= "Complementarity determining region"
                                                                                                                                     'label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                               label=
                                                                                                                                                                                                                            'label= Synthetic peptide linker
note= "Links the heavy chain to
                                                                                                                                                                                                                                                                                                'label= CDR
'note= "Complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                               note= "Complementarity
                                                                                                                                                                                                                                                                                                                                                                                                      .abel= CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
                                                                                                                                                                                                                                                                                .152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                    .249
                                                                                                                                                                                                       .186
                                                                                                                                                                                                                                                                                                                                                          .126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     87pp; English.
                                                                                                                             .208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 97;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                         determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 246;
                                                                                                                                                                                                                              the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                   region"
                                                                                                                                                                                                                         light chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunoconjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for
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RESULT 8
AAU04945
ID AAU0
XX
AC AAU0
XX
DT 06-A
DT 24-0
XX
DE Huma
XX
Huma
XX
Homo
OS Mus
OS Mus
Synt
OS Chim
XX
FT Pept
FT Prot
FT Prot
FT Prot
FT Regi
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                                                                                                                                                                                                                                                                                                                                                                                                               Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                               The sequence represents a humanised anti-p185 single chain antibody which is linked to a human interleukin-2 (II-2) molecule to make a fusion protein. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cancers particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, renal system, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
Region
                                                                                                           Mus sp.
Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel immunoconjugate comprises a humanized
                     Protein
                                          Protein
                                                               Peptide
                                                                                                  Chimeric.
                                                                                                                                   Homo
                                                                                                                                                       cancer;
                                                                                                                                                                 Humanised;
                                                                                                                                                                                      Humanised
                                                                                                                                                                                                           06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                             AAU04945;
                                                                                                                                                                                                                                                                  AAU04945 standard; protein; 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 7; Fig 9; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Austin R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JAN-2000; 2000US-0177258P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JAN-2001; 2001WO-US001919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CHIR ) CHIRON CORP
(HAMI-) HAMILTON CIV
(HAMI-) HAMILTON REC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200153354-A2
                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-451904/48.
                                                                                                                                                                                                                                                                                                                        72
                                                                                                                                                                                                                                                                                                                                             1 WINTYTGOSTYADDFKE
                                                                                                                                                                                                                                                                                                                                                                  Similarity 17; Conserv
                                                                                                                                                       tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS09507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HAMILTON CIVIC HOS
                                                                                                                                                                                                                                                                                                                       WINTYTGOSTYADDFKE
                                                                                                                                                                                     anti-p185 antibody/IL-2 fusion protein.
                                                                                                                                                               antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                           (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                               AA;
/label= Mature_fusion_protein
23. .259
/label= Humanised_antibody_520
53. .57
                                          /label= Signal_peptide
23. .409
                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     260. .267
/label= Glu_Glu_epitope
                                                                                                                                                                                                            entry)
                                                                                                                                                     p185; 520C9H; interleukin-2; ocarcinoma; fusion protein.
                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for inhibiting tumor cell growth in vivo anti-p185 antibody linked to an Interleukin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HOSPITALS
         Humanised_antibody_520C9H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BB;
                                                                                                                                                                                                                                                                                                                        88
                                                                                                                                                                                                                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                              Score 97;
Pred. No.
                                                                                                                                                                                                                                                                   ₽
                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RES DEV INC.
CENT.
                                                                                                                                                                                                                                                                                                                                                                   1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                         DB 4;
                                                                                                                                                                                                                                                                                                                                                                                      Length 267;
                                                                                                                                                                IL-2;
                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                immunoconjugate;
                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                   Gaps
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문
                 5
                                                                                     Query Match
Best Local S
Matches 17
                                                                                                         The sequence represents a humanised anti-p185 (520C9) single chain antibody/human interleukin-2 (IL-2) fusion protein. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cances particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, rensestem, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                Claim 13; Fig 11; 74pp; English.
                                                                                                                                                                                                                                 Novel immunoconjugate comprises a humanized
                                                                                     Sequence
                                                                                                                                                                                                                                                                    N-PSDB; AAS09508.
                                                                                                                                                                                                                                                                               WPI; 2001-451904/48.
                                                                                                                                                                                                                                                                                                      Austin R,
                                                                                                                                                                                                                                                                                                                                      (CHIR )
(HAMI-)
                                                                                                                                                                                                                                                                                                                                                                     20-JAN-2000; 2000US-0177258P
                                                                                                                                                                                                                                                                                                                                                                                            19-JAN-2001; 2001WO-US001919
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Region
                                                                                                                                                                                                                                                                                                                            (HAMI-)
                                         y Match 100.0%;
Local Similarity 100.0%;
hes 17; Conservative 0;
72
                   μ
                                                                                                                                                                                                                                                                                                                         CHIRON CORP.
HAMILTON CIVIC HOSPITALS
HAMILTON REGIONAL CANCER
                   WINTYTGOSTYADDFKE 17
WINTYTGOSTYADDFKE
                                                                                     409 AA;
                                                                                                                                                                                                                                                                                                      Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         406
                                                                                                                                                                                                                                                                                                                                                                                                                                                             note= "Encode by ACA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= Synthetic_linker_peptide
note= "Links the antibody to the IL-2 molecule"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note= "Complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= CDR
/note= "Complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= CDR
/note= "Complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label= CDR
/note= "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label= Synthetic_peptide_linker
/note= "Links the heavy chain to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  notes "Complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              label= CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "Encoded by GGG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .208
                                                                                                                                                                                                                                                                                                    Ring DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 186
                                                                                                                                                                                                                                  useful for inhibiting tumor cell growth in vivo anti-pl85 antibody linked to an Interleukin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IL_2
88
                                         Score 97; DB 4;
Pred. No. 1.9e-06;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                            RES DEV INC.
CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        determining region"
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                                                              Length 409;
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                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        region'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  region"
                                         0
                                         Gaps
                                                                                                                                                     cancers
                                                                                                                               renal
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RESULT 9

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AAR39571
ID AAR3
XX AAR3
XX AAR3
XX AAR3
XX Seq.
XX Seq.
XX Synt
XX WO9:
XX WO9:
XX WO9:
XX WO9:
XX WO9:
XX WO9:
XX WPI
DR N-P;
XX WPI
DR N-P;
XX WPI
DR N-P;
XX WPI
CC C-e
CC C-e
CC CO1
CC Si3
CC fie
XX Seq
RESULT 10
AAB28159
ID AAB28
XX AAB28
XX AAB28
XX AAB28
DT 08-FE
XX Murin
XX Murin
XX C-erb
XX Mus 8
XX Mus 9
PN W0200
XX Mus 8
XX Mus 9
PN W0200
XX TO APP
XX TO APP
XX TO APP
XX TO AB28
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                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 17
           10-APR-2000; 2000WO-EP003176
                                                                                                           Murine; immunoglobulin; antigen-binding; framework region; carcinoma; c-erbB2; carcinoma-associated antigen.
                                                                                                                                                                                                                      AAB28159 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 534 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                      c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. Such as breast and ovarian tumour cells, which is an approx. 200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New single chain Fv polypeptide binding to C-erbB-2 tumour antigen - imaging or treating breast or ovarian cancer etc.
                                     19-OCT-2000
                                                            WO200061635-A2
                                                                                    da snw
                                                                                                                                              Murine anti-EGP-2 single chain Fv fragment.
                                                                                                                                                                        08-FEB-2001
                                                                                                                                                                                                AAB28159;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1993-272889/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Huston JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-FEB-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumour antigen; c-erbB-2; G-FIT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
07-FEB-1994
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ46086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 65-68; 87pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Houston
                                                                                                                                                                                                                                                                                                                                  100.0%; llarity 100.0%; Conservative 0;
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(first en
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Pred. No. 2.5e-06;
0; Mismatches 0;
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UNIV ZUERICH PLUECKTHUN A.

99EP-00107030

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RESULT 11
AAW35131
ID AAW35
XX AW35
XX 20-AP
XX RNase
KW RNase
KW tumou
XX Synth
XX W0973
XX 28-AU
PF 19-FE
XX 21-FE
XX 21-FE
XX W1;
DR W1-FE
XX W1;
DR W1-FSI
XX WP1;
DR W1-PSI
XX W1,
XX Rana
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunoglobulin (Ig). The method comprises identifying antigen-binding groups derived from donor Ig and framework regions derived from an acceptor Ig. The present sequence is one such donor Ig fragment. One or more of the residues present at the positions in the present sequence are replaced by those present at the corresponding positions in the donor Ig, after comparing the structural features of the VH domains of the acceptor Ig and the donor Ig. The acceptor Ig fragment used in the present invention is human anti-c-Erbb2 single chain Fv fragment 4D5 (AAB28158). The method of the present invention is useful for producing a pharmaceutical composition which can be useful for treating human and the present invention be useful for treating human and the present invention be useful for treating human and the present invention be useful for treating human and the present invention be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for treating human and the composition which can be useful for the composition which can be useful for the composition and the composition which can be useful for the composition and the composition which can be useful for the composition and the composition which can be useful for the composition and the composition and the composition and the composition and the composition a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rana pipiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6; Page 51; 51pp; English.
                                                                                              WPI; 1997-435168/40.
N-PSDB; AAT94969.
                                                                                                                                                                                                                                                                                                              19-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                              28-AUG-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R. pipiens recombinant RNase rOnc fusion protein 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW35131;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for stabilising a chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-679468/66.
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                        particularly tumour cells.
                                                 Ribonuclease molecules based on native Onconase
                                                                                                                                                                      Rybak SM,
                                                                                                                                                                                                                                                                  21-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                             WO9731116-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNase A; ribonuclease; cytotoxic; tumour cell growth; frog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           carcinomas, since c-erbB2 is a carcinoma-associated antigen
                                                                                                                                                                                                                     (USSH ) US
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                                                                                                                                                                      Newton DL,
                                                                                                                                                                                                                     DEPT HEALTH & HUMAN SERVICES.
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93.8%;
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Pred. No. 1.9e
1; Mismatches
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                                                                                                                                                                      Wlodawer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nOnc; immunofusion;
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                                                 used for killing
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RESULT 12
AAW35132
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Matches
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Sequences AAW35125 to AAW35135 represent recombinant fusion proteins (rOnc) which are modifications of the RNase Onconase (RTM) (nOnc). Such novel ribonuclease molecules are highly cytotoxic and can be used alone or to form chemical conjugates or to target recombinant immunofusions. They are used particularly for decreasing tumour cell growth. They can also be used for cell separation in vitro by selectively killing unwanted types of cells, e.g. in bone marrow prior to transplantation into a patient undergoing marrow ablation by radiation, or for killing leukaemia cells or T-cells that would cause graft versus host disease. The toxins can also be used to selectively kill unwanted cells in culture. The new ribonucleases have increased cytotoxic activity compared to nonc and also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNase A; ribonuclease; cytotoxic; onconase; nOnc; immunofusion; tumour cell growth; frog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R. pipiens recombinant RNase rOnc fusion protein 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW35132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW35132 standard; protein; 366 AA
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                                                                                                                                                                                                                                                   Ribonuclease molecules based on native particularly tumour cells.
                                                                                                                                                                                                                                                                                                           N-PSDB; AAT94970
                                                                                                                                                                                                                                                                                                                                                             Rybak SM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                   21-FEB-1996;
                                                                                                                                                                                                                   Disclosure; Page
                                                                                                                                                                                                                                                                                                                             WPI; 1997-435168/40
                                                                                                                                                                                                                                                                                                                                                                                                (USSH ) US DEPT HEALTH & HUMAN SERVICES
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                                                                                                                                                                                                                   74; 90pp;
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93.8%;
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                                                                                                                                                                                                                     English
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Pred. No. 2.9e-05;
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                                                                                                                                                                                                                                                                         Onconase - used
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                                                                                                                                                                                                                                                                           for killing cells,
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RESULT 13
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AAP70625
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AC AAP70
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Best Local
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                                             (ITGE-)
(ROBI/)
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09-APR-1991
                Robinson RR,
                                                                                          01-NOV-1985;
                                                                                                                                                         07-MAY-1987.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence encoded by the V region of the L6 VH cDNA clone pH3-6a.
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                                                                                                                         27-OCT-1986;
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                                             INT GENETIC ENG INC. ROBINSON R R.
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(first entry)
                Liu AY, Horwitz AH,
                                                                                          85US-00793980
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/*tag= __
/label= FR4
/40, .183
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/label= JH2
129. .139
/*tag= i
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/label= FR3
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/label= CDR2
86. .117
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label= CDR1
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Pred. No. 2.
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              Wall R;
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AAP82936
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03-OCT-2002
12-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The patentors claim a chimeric antibody molecule comprising 2 light chains and 2 heavy chains, each comprising a constant human region and a variable non-human region. Coding sequences for the Ig chains are also claimed. The invention provides consensus sequences of light and heavy chain J regions useful in the design of oligonucleotides (UIGs) for use as primers or probes for cloning immunoglobulinlight or heavy chain mRNAs or genes. Depending on the nature of design of a particular UIG, it may be capable of hybridizing to all Ig mRNAs or genes containing a single specific J sequence. UIG denotes universal immunoglobulin gene. (Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                       Key
                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse immunoglobulin; variable region; tumour antigen; chimeric antibodies; L6Vh; pH3-6a.
 05-MAY-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP82936 standard; protein; 183 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 183 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antigens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               procedures, with chimeric antibodies etc. related to cancer specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; AAN70969.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ch 89.7%;
l Similarity 93.8%;
15; Conservative
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(revised)
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/label= Jh2
129. .183
/label= FR4
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/label= CDR3
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6. .117
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Pred. No. 2.8e-05;
0; Mismatches 1
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A library primed by oligo (dT) on L6 poly (A+) RNA was screened for kappa clones with a mouse Ck region probe. From the L6 library, several clones were isolated. A second screen with a 5' JK5 specific probe identified the L6 (JK5) light-chain clones. Heavy chain clones of L6 were isolated be screening with the Jh2 oligonucleotide. The heavy and light chain genes from the clones pH3-6a (and pL3-12a) were inserted into M13 bacteriophage vectors of which the sequence was determined by the dideoxytermination method. Amino acid residues 26-35 have been confirmed by peptide sequence. See also AAN82415-20. (Updated on 03-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ONCO )
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N-PSDB; AAN82417.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page ?; 83pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polynucleotide encoding variable region of mouse immunoglobulin - specific for human tumour antigen, and expression products, including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Liu
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03-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                immunoglobulin; variable region; tumour antigen;
ric antibodies; Bal-31 deletion clone M13mp19-C1-delta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric human-mouse antibodies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WINTYTGOPTYADDFK 84
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     86. .117
/label= FR3
118. .128
                                                                                                                     50. .54
/label= CDR1
55. .68
/label= FR2
                                                                                                                                                                                                                                                /label= leader_peptide
20. .49
                                                                           /label= CDR2
                                                                                                                                                                                                               /label= FR1
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                                                                                                    . 85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hellstrom
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Pred. No.
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2.8e-05;
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Matches 15; Conserv
                                                                                                                                                 The nuclease Bal-31 was used to chew away the oligo (dC) in the L6 Vh clone pH3-6a (AAN82417). For this clone nucleotides 1-38 were deleted. Then the sequence was modified by site-directed mutagenesis with the MJF ApaI primer. This 31-base primer introduces an ApaI site in the mouse of gene at position at a position analogous to human Cgammal cDNA gene module. See also AAN82415-20. (Updated on 03-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                            New polynucleotide encoding variable region of mouse immunoglobulin - specific for human tumour antigen, and expression products, including complete chimeric human-mouse antibodies.
                                                                                                                                                                                                                                                                                                                    WPI;
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                                                                                                                              Sequence 183
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DB; AAN82419.
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125. .139
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93.8%;
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2.8e-05;
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                                                                                                                                                                                                                               Score
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1: /cgn2=6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

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Copyright (c) 1993 - 2005 Compugen Ltd.
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   US-09-887-853-6
7 US-10-683-547-6
US-09-766-543-10
US-09-766-543-12
US-09-971-543-8
US-09-971-485-102
4 US-10-071-485-91
4 US-10-071-485-91
5 US-10-071-485-91
6 US-10-071-485-91
7 US-10-071-485-91
8 US-10-071-485-91
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Sequence 6, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 12, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 102, Appli
Sequence 93, Appli
Sequence 91, Appli
Sequence 67, Appli
Sequence 67, Appli
Sequence 85, Appli
Sequence 85, Appli
Sequence 89, Appli
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ALIGNMENTS

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US-09-887-853-6

Sequence 6, application US/09887853

Patent No. US2020168375A1

GENERAL INFORMATION:

APPLICANT: Huston, James S.

Oppermann, Hermann
Houston, L. L.

Ring, David B.

RITLE OF INVENTION: Biosynthetic Binding Proteins For Imaging

NUMBER OF SEQUENCES: 11

CORRESPONDENCE ADDRESS:

ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department STREET: Exchange Place, 53 State Street
CITY: Boston
STATE: Massachusetts
COUNTRY: USA
ZIP: 02109

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25

PAPLICATION NUMBER: US/09/887,853
FILING DATE: 21-Un-2001
CLASSIFICATION NUMBER: US/09/887,853
FILING DATE: 21-Un-2001
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/887,853
FILING DATE: CUNKnown>
PRIOR APPLICATION NUMBER: US/08/133,804
FILING DATE: CUNKnown>
ATTORNEY/AGENT INFORMATION:

NAME: Kelley, Robin D.
REGISTRATION NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
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TELEFAX: 617-248-7100 INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:

TELEPHONE:

617-248-7477

0

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; Sequence 10, Application US/09766543
; Patent No. US20020041865A1
; GENERAL INFORMATION:
; APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Ring, David B.
; APPLICANT: Ring, David B.
; TITLE OF INVENTION: METHODS FOR TREATING TUMORS
; FILE REFERENCE: PP01679.002
; CURRENT FILING DATE: 2000-01-19
; CURRENT FILING DATE: 2000-01-19
; CURRENT FILING DATE: 2000-01-19
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HOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-887-853-6
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Best Local S
Matches 17
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 6
LENGTH: 243
TYPE: PRT
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Publication No. US20050058638A1
GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING
FILE REFERENCE: CIET-P01-130

CURRENT APPLICATION NUMBER: US/10/683,547

CURRENT FILING DATE: 2003-10-10

PRIOR APPLICATION NUMBER: US/09/558,741

PRIOR APPLICATION NUMBER: US/09/558,741

PRIOR PILING DATE: 2000-04-26

PRIOR PILING DATE: 2000-04-26

PRIOR FILING DATE: 1992-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Huston, J. APPLICANT: Houston, L.L. APPLICANT: Ring, D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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Pred. No.
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PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOPTWARE: PATENTIN VEY. 2.0
SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Ring, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PP01679.002
                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/09971543
Patent No. US20020146846A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12, Application US/09766543
Patent No. US20020041865A1
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PRIOR APPLICATION NUMBER: PCT/EP00/03176
PRIOR FILING DATE: 2000-04-10
PRIOR APPLICATION NUMBER: EP 99 10 7030.1
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
                                                                                                                  APPLICANT: PLUCKTHUN, ANDREAS
APPLICANT: HONEGGER, ANNEMARIE
APPLICANT: WILLUDA, JORG
TITLE OF INVENTION: NOVEL METHOD FOR THE STABILIZATION OF CHIMERIC
TITLE OF INVENTION: IMMUNOGLOBULINS OR IMMUNOGLOBULIN FRAGMENTS, AND
TITLE OF INVENTION: STABILIZED ANTI-EGP-2 SCFV FRAGMENT
FILE REFERENCE: PLUCK-3 CON
CURRENT APPLICATION NUMBER: US/09/971,543
CURRENT FILING DATE: 2001-10-04
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CURRENT FILING DATE: 2000-01-19
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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17; Conserv
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17; Conserv
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Pred. No. 3.6e-07;
; Mismatches 0;
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Pred. No. 3.8e-07;
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; ORGANISM: Mus sp. US-09-971-543-8
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Sequence 102, App
Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:

APPLICANT: PLUCKTHUN, ANDREAS
APPLICANT: HONEGGER, ANNEMARIE
APPLICANT: HONEGGER, ANNEMARIE
APPLICANT: WILLUDA, JORG

TITLE OF INVENTION: NOVEL METHOD FOR THE STABILIZATION OF CHIMERIC
TITLE OF INVENTION: IMMUNOSLOBULINS OR IMMUNOGLOBULIN FRAGMENTS, A
TITLE OF INVENTION: STABILIZED ANTI-EGP-2 SCFV FRAGMENT
FILE REFERENCE: PLUCK-3 CON
CURRENT APPLICATION NUMBER: US/09/971,543
CURRENT FILING DATE: 2001-10-04
PRIOR APPLICATION NUMBER: PCT/ED00/03176
PRIOR APPLICATION NUMBER: EP 99 10 7030.1
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LENGTH: 253
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Patent No. US20020146846A1
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                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                   PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-08-14
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
                                                                                                                                                    TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC TITLE OF INVENTION: SHOCK, TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISCRIBERS FILE REFERENCE: INNS:015
CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 09/485,737
PRIOR FILING DATE: 2000-02-14
                    NUMBER OF SEQ ID NOS: 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 116
TYPE: PRT
                                                                                                                                                                                                                                                                                                                 PPLICANT: Buyse, Marie-Ange
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Mus sp.
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PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                           Application US/10071485
p. US20030099648A1
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93.8%;
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93.8%;
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Pred. No.
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Pred. No.
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hes 0; Indels
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US-10-071-485-91
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                                                                                                                                                                                                                                                      Sequence 91, Ap
Publication No.
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Best Local Similarity
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Best Local Similarity
APPLICANT: Buyse, Marie-Ange
APPLICANT: Sablon, Erwin
APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 09/485,737
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 2000-02-14
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-08-14
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TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/485,737
PRIOR FILING DATE: 2000-02-14
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-08-14
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR APPLICATION NUMBER: EPO 97870122.5
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CURRENT FILING DATE: 2002-02-07
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: SYNTHETIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 230
TYPE: PRT
ORGANISM: Artificial Sequence
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Pred. No. 2.5e
1; Mismatches
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2.5e-05;
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US-10-071-485-67
; Sequence 67, Application US/10071485
; Publication No. US20030099648A1
; GENERAL INFORMATION:
; APPLICANT: Buyse, Marie-Ange
; APPLICANT: Sablon, Erwin
; TITLE OF INVENTION: INTERFERON-gamma-I
; TITLE OF INVENTION: CACHEXIA, IMMUNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 09/485,737
PRIOR FILING DATE: 2000-02-14
PRIOR FILING DATE: 2000-02-14
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR FILING DATE: 1998-06-18
PRIOR FILING DATE: 1998-06-18
PRIOR PRIOR DATE: 1998-06-18
PRIOR PRIOR DATE: 1998-06-18
PRIOR PRIOR DATE: 1998-06-19
PRIOR PRIOR DATE: 1998-06-19
PRIOR PRIOR DATE: 1998-06-19
PRIOR PRIOR DATE: 1998-06-19
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US-10-071-485-2
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; OTHER INFORMATION: SYNTHETIC
US-10-071-485-91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1997-08-18
NUMBER OF SEQ ID NOS: 104
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2
LENGTH: 267
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Publication No. US20030099648A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
NUMBER OF SEQ ID NOS: 104
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Synthetic
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                Conservative
                               INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC SHOCK,
CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
                                                                                                                                                                                                                                                                                                                                                                  87.6%;
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87.5%;
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Pred. No. 2.9e.
1; Mismatches
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Pred. No. 2.
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2.9e-05;
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2.6e-05;
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                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 09/485,737
PRIOR FILING DATE: 2000-02-14
PRIOR PEPLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-08-14
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
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PRIOR FILING DATE: 2000-02-14
PRIOR PEDLICATION NUMBER: PCT/EP 98/05165
PRIOR FILING DATE: 1998-08-14
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR FILING DATE: 1997-08-18
PRIOR FILING DATE: 1997-08-18
                                                                            Matches
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Matches
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                                                                                                                Query Match
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Publication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Buyse, Marie-Ange APPLICANT: Sablon, Erwin
                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence
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CURRENT FILING DATE: 2002-02-07
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                                                                                                                                                                        OTHER INFORMATION: SYNTHETIC
                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: SYNTHETIC
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                                                                                              Local
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50 WINTYTGESTYVDDFK 65
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                                  1 WINTYTGQSTYADDFK 16
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                                                                                            Similarity
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Similarity 87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/10071485
10. US20030099648A1
                                                                          Conservative
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                                                                                            87.6%;
                                                                                            Score 85;
Pred. No.
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Pred. No. 5.1e-05;
                                                                          Mismatches
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                                                                                                                DB 14;
                                                                                            9e-05;
                                                                                                            Length 541;
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RESULT 13 US-10-071-485-90 ; Sequence 90, Application US/10071485

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CURRENT APPLICATION NUMBER: US/09/791,551
CURRENT FILING DATE: 2001-02-26
PRIOR APPLICATION NUMBER: 60/15,390
PRIOR FILING DATE: 2000-02-28
PRIOR APPLICATION NUMBER: 60/233,625
PRIOR APPLICATION NUMBER: 60/233,625
PRIOR FILING DATE: 2000-09-18
NUMBER OF SEQ ID NOS: 119
SOFTMARE: Patentin Ver. 2.1
SEQ ID NO 86
LENGTH: 17
TYPE: PRT
ORGANISM: Mus sp.
US-09-791-551-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 90
LENGTH: 711
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: SYNTHETIC
US-10-071-485-90
RESULT 15
US-10-018-245A-2
### Sequence 2, Application US/10018245A
#### Publication No. US20040115196A1
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Sequence 86, Application US/09791551

Publication No. US20030235584A1

GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-02-14
PRIOR PPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR PILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: EPO 97870122.5
PRIOR PILING DATE: 1997-08-18
PRIOR PILING DATE: 1997-08-18
NUMBER OF SEQ ID NOS: 104
                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                            Query Match
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KLOETZER, WILLIAM S.
APPLICANT: HANNA, NABIL
TITLE OF INVENTION: METHOD FOR PREPARING ANTI-MIF ANTIBODIES
FILE REFERENCE: 037003/0277869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sablon, Erwin
TITLE OP INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING
TITLE OP INVENTION: SHOCK,
TITLE OP INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS:015
CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
CURRENT FILING DATE: 2002-02-07
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                                                                                                                                                                                                                                                   14;
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Similarity 87.5%; Pred. No. 2.4e-06;
14; Conservative 1; Mismatches 1; Indels
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Search completed: April Job time : 26.0927 secs
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                                                                                                                                                                                                                            ; TYPE: PRT
ORGANISM: mouse
; PEATURE:
; CTHER INFORMATION: CDR-H2 of anti-human TNF-alpha antibody
US-10-018-245A-2
                                                                                                                                                     Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/018,245A
CURRENT FILING DATE: 2002-07-12
PRIOR APPLICATION NUMBER: JP 117394/2000
PRIOR FILING DATE: 2000-04-19
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 2
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Novel recombinant antibody, amino acid sequences of its complement TITLE OF INVENTION: determining regions and genes encoding the same FILE REFERENCE: 46224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: NAGAHIRA, Kazuhiro
APPLICANT: NAKANISHI, Toshihiro
                                                                                                                                                cn 86.6%;
l Similarity 87.5%;
l4; Conservative
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                                                                          WINTYTGEPTYADDFK 16
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1; Mismatches
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         9
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(without alignments)
266.512 Million cell updates/sec
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LG nAb heavy chain pre
BHD9D10 protein -
Ig heavy chain V r
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A;Molecule type: mRNA
A;Residues: 1-115 <WEI>
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Matches 15
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ALIGNMENTS

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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-120 <FE2>
A;Crose-references: EMBL:M90691
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                          A;Status: FILE TYPE: DNA A;Rolecule type: DNA A;Rolecule type: DNA A;Rolecule: 1-120 <FEL>
A;Residues: 1-120 <FEL>
A;Residues: 1-20 <FEL>
A;Cross-references: GB:M90690; NID:g195065; PIDN:AAA38146.1; PID:g195066
A;Cross-references: GB:M90690; NID:g195065; PIDN:AAA38146.1; PID:g195066
                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 2
B42848
L6 mAb heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Apate: 27-Apx-1993 #sequence revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: B42848; S33903
R;Fell, H.P.; Gayle, M.A.; Yelton, D.; Lipsich, L.; Schieven, G.L.; Marken, J.S.; Aruffo, J. Biol. Chem. 267, 15552-15558, 1992
A;Title: Chimeric L6 anti-tumor antibody. Genomic construction, expression, and character A;Reference number: A42848; MUID:92348410; PMID:1639794
A;Accession: B42848; MUID:92348410; PMID:1639794
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C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 09-Jul-2004
C;Accession: S19968
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7.4e-07;
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A;Reference number: S35759
A;Accession: S35759
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-136 <FRO>
A;Accessation type: DNA
A;Molecule type: DNA
A;Residues: 1-101 <WIN>
A;Residues: 1-101 <WIN>
A;Cross-references: GB:X03301; NID:g51757; PIDN:CAA27040.1; PID:g773215
A;Cross-references was determined from the differentiated gene
                                                                                                         Ig heavy chain V region (VGAM3-8) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Nov-1988 #sequence_revision 19-Nov-1988 #text_change 23-Jul-1999
C;Accession: D24672
R;Winter, E.; Radbruch, A.; Krawinkel, U.
EMBO J. 4, 2861-2867, 1985
A;Reference number: A91022; MUID:86055722; PMID:2998759
A;Accession: D24672
                                                                                                                                                                                                                                                                                                                                           문
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BHD9D10 protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 21-Jan-2000
C;Accession: S35759
R;Froyen, G.F.V.
submitted to the EMBL Data Library, May 1993
A;Reference number: S35759
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S35759
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A;Title: Chimeric mouse-human IgG1 antibody that can mediate lysis of cancer cells. A;Reference number: S33905; MUID:87204152; PMID:3106970
A;Accession: S33905
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A; Residues: 1-146 <LIU>
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                         WINTYTGESTYVDDFK 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WINTYTGOPTYADDFK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WINTYTGOPTYADDFK 65
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                  87.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          89.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  89.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 85; DB 2;
Pred. No. 1.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 87; DB 4;
Pred. No. 9.6e-07;
0; Mismatches 1
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Pred. No. 7.8e-07;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 136;
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                                                                 submitted to the EMBL Data Library, A;Reference number: S24776 A;Accession: S24778
                                             A; Molecule type: DNA
                                                                                                                                                                                                                                                                                     R;Klages, S. submitted to the EMBL Data Library, A;Reference number: $24763
                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
S24765
                                                                                                                          R;Thomas, J.W.
                                                                                                                                                                                                A;Cross-references: EMBL:Z15001
A;Accession: S24773
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                                                                                                                                        A;Cross-references: EMBL:Z15013
                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-105 < KLW>
                                                                                                                                                                                                                                                     A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Thomas, J.W. submitted to the EMBL Data Library,
                                                                                                                                                                                                                                     A;Residues: 1-105 <KLA>
                                                                                                                                                                                                                                                                           A;Accession: S24765
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A; Residues: 1-105 < THO>
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A;Accession: S24777
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A; Residues: 1-105 < KLA>
A; Cross-references: EMB
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S24764
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A; Residues: 1-105 < KLW >
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Reference number: S24763
A; Accession: S24764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 submitted to the EMBL Data Library, August 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Klages,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 86.6%;
Similarity 87.5%;
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WINTYTGEPTYADDFK 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WINTYTGQSTYADDFK 16
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Pred. No. 2.1e-06;
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Pred. No. 2e-06;
1; Mismatches
                                                                                                     August 1992
                                                                                                                                                                                                                                                                                                               August 1992
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Ig heavy chain V region (subgroup XI) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Apecies: Mus musculus (house mouse)
C;Apecies: Peb-1995 #sequence revision 21-Nov-1998 #text_change 23-Jul-1999
C;Accession: S24765; S24773; S24778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: EMBL:Z15020; NID:g52616; PIDN:CAA78739.1; PID:g52617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig heavy chain V region (subgroup XI) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 20-Feb-1995 #sequence revision 21-Nov-1998 #text_change 23-Jul-1999
C;Accession: S24764; S24772; S24777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Superfamily: immunoglobulin V region; immunoglobulin homology; Keywords: heterotetramer; immunoglobulin
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Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C;Accession: $26325
R;Stark, S.E.; Caton, A.J.
J. Exp. Med. 174, 613-624, 1991
A;Title: Antibodies that are specific for a single amino acid interchange in a protein A;Accession: $26309; MUID:91341421; PMID:1908510
A;Accession: $26325
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-109 <STMA>
A;Residues: 1-109 <STMA>
A;Residues: 1-109 <STMA>
A;Cross-references: UNIPROT:Q921A6; EMBL:X59210; NID:g52080; PID:g1334043
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Kcywords: heterotetramer; immunoglobulin homology <TMMA>
                                                                                                                                                                                                                                                                                                                                                             Ig heavy chain V region (M-T406) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 09-Jul-2004
C;Accession: S19967
R;Weissenhorn, W.; Riethmueller, G.; Weiss, E.M.; Rieber, E.P.
aubmitted to the EMBL Data Library, March 1992
A;Description: Structural characterization of CD4 mAb.
A;Reference number: S19963
A;Accession: S19963
A;Accession: S19963
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S19967
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A;Introns: 9/1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Koywords: heterotetramer; immunoglobulin
RESULT 10
A53285
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                                                                                                                                                                                                                                 ;12-95/Domain:
                                                                                                                                                                                                                                                                                                           ;Molecule type: mRNA
;Residues: 1-118 <WEI>
                                                                                                                                                                                                                                                                                                                                             ;Status: preliminary
                                                                                                                                                          Matches
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Best Local (
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Best Local
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Best Local
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                                                                              47
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                                                                                                                1 WINTYTGOSTYADDFK 16
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                                                                                                                                                                          Similarity
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                                                                            WINTYTGEPTYADDFK 62
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                                                                                                                                                                                                                                 immunoglobulin homology <IMM>
                                                                                                                                                      Conservative
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                                                                                                                                                                        86.6%;
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                                                                                                                                                                        Score 84; DB 2;
Pred. No. 2.3e-0
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                                                                                                                                                                                                                               RESULT 12
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                  A;Status: preliminary
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Best Local
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Best Local :
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C32530
C32530
Ig heavy chain V region (10/8) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 03-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 16-Aug-1996
C;Accession: C32530
R;Deverson, B:, Berek, C:, Taussig, M:, Feinstein, A.
Eur. J. Immunol. 17, 9-13, 1987
By Title: Monoclonal BALB/C anti-progesterone antibodies use family IX varial A;Accession: C32530
A;Accession: C32530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Introns: 16/1
C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
F;35-118/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PH1404
If heavy chain V region (clone micro m+ 46-12) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 21-Jan-2000
C;Accession: PH1404; PH1406
C;Accession: PH1404; PH1406
R;Shirasawa, T.; Miyazoe, I.; Hagiwara, S.; Kimoto, H.; Shigemoto, K.; Taniguchi, M.; Taly. Exp. Med. 176, 1209-1214, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Accession: PH1406
A;Molecule type: DNA
A;Residues: 115-121 <SH2>
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A;Title: Molecular characterization of monoclonal anti-steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig heavy chain V and J regions, monoclonal antibody SCET.M8.1 - mouse (fragment) c;Species: Mus musculus (house mouse) C;Decies: Mus musculus (house mouse) C;Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 20-Jun-2000 C;Accession: A53285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-124 <SHI>
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A;Molecule type: DNA; protein
A;Residues: 1-119 <SAW>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: PH1403; MUID: 93018837; PMID: 1402663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Title: Heavy chain variable (VH) region diversity generated by
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2.5e-06;
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Ly neavy chain V region TE32 - mouse
C;Species: Mus musculus (house mouse)
C;Accession: D32967
C;Accession: D32967
R;Levy, R: Assulin, O.; Scherf, T.; Levitt, M.; Anglister, J.
Biochemistry 28, 7168-7175, 1989
A;Title: Probing antibody diversity by 2D NMR: comparison of amino acid sequences, predictive: probing antibody diversity by 2D NMR: comparison of amino acid sequences, predictive: predictive: name and acid sequence predictive: prediction: D32967
A;Status: preliminary; nucleic acid sequence not shown; not compared with conceptual tradictive: predictive: manuacylobulin V region; immunoglobulin V region; immun
RESULT 15
C32967
Ig heavy chain V region TE33 - mouse
C;Species: Mus musculus (house mouse)
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C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 09-Jul-2004
C;Accession: S19965
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A;Residues: 1-102 <DEV>
A;Cross-references: GB:M27585
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A; Description: Structural characterization of CD4 mAb.
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;8-91/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-115 <WEI>
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A; Status: preliminary
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                                                                                                                                                                                                                              WINTYSGVPTYADDFK 65
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llarity 81.2%;
Conservative
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Pred. No. 4.4e-05;
1; Mismatches 2
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Pred. No. 8.9e-06;
1; Mismatches 2;
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C;Date: 29-Jan-1990 #sequence_revision 29-Jan-1990 #text_change 21-Jan-2000 C;Accession: C32967 R;Levy, R.; Assulin, O.; Scherf, T.; Levitt, M.; Anglister, J. Biochemistry 28, 7168-7175, 1989 A;Title: Probing antibody diversity by 2D NMR: comparison of amino acid sequence number: A32967; MUID:90057406; PMID:2819059 A;Accession: C32967
                                                                                                                                                                                                                                               C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
Search completed: April 4, 2005, 15:58:00 Job time: 6.13738 secs
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A; Residues: 1-114 < LEV >
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                                                                                                                                                                                         Query Match
Best Local :
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                                                                                                                                                                                         Similarity
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81.2%;
                                                                                                                                                                     Score 76; DB 2;
Pred. No. 4.4e-05
1; Mismatches
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Maximum Match 100%
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: uniprot_sprot:*
2: uniprot_trembl:*
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  GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Q9VTS5
Q7REI4
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Q9JL79
Q65ZL3
Q683Y7
Q683Y8
Q99LA6
Q6PIL0
Q925S1
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Q7N848
Q7YTW2
Q6N091
Q6A3Y7
Q8A3Y7
Q7RXD7
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Q9NA29
Q8YTW6
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Q92:a6
Q92:a7
Q65:13
Q683Y7
Q6813Y8
Q991a6
Q901a6
Q60:10
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1 lactococcus
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4 plasmodium
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8 photorhabdu
2 caenorhabdi
1 homo sapien
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ULT 2 29JL: Q9JL: Q9JL: Q1-O0 01-O0 01-O0 Anti: (Frag Mus r Eukal		Query Mat Best Loca Matches	PIR; PIR; PIR; PIR; PROS; NON NON SEQUI	SEQUI MEDL: Chung Yi K "Clo monoo gene: Mol.	Mus t Euka: Mamma NCBI	O1-DI O1-DI Anti	SULT 1 21A6 0921		44444 0) A B B B B B B B B B B B B B B B B B B	3
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ee) egion a, Eutoleostomi;		Length 241; ; Indels 0; Gaps 0;	CRC64;	I.H., Lee S.D., VH and VL of a tive with NCA-95 and	a; Euteleostomi; ae; Murinae; Mus.	(e)			404040	Often active coact Often close tridium Ofvuh9 drosophila Ofkul0 drosophila Often drosophila	7

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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

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InterPro; IPR007110; Ig-like.
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SMART; SM00409; IG; 1.
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Q65ZL3;
25-OCT-2004
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MEDLINE-96269537; PubMed-8699031; DOI=10.1016/0022-1759(96)00043-9;

NOSEL D., Bernardi T., Navarro-Teulon I., Marin M., Martinetto J.P.,

Ducancel F., Mani J.C., Pau B., Piechaczyk M., Biard-Piechaczyk M.;

Place of the individual contributions of immunoglobulin heavy and the chains to the binding of antigen using cell transfection and light chains to the binding of antigen using cell transfection and
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J. Immunol. Methods 193:177-187(1996).
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Mammalia; Eutheria;
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EMBL; AF206027; AAF69325.1; -.
HSSP; P01751; INQB.
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Malkiel S., Liao L., Cunningham M.W., Diamond B.;
MT-Cell-dependent antibody response to the dominant epitope of
streptococcal polysaccharide, N-acetyl-glucosamine, is cross-reactive
with cardiac myosin.";
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InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
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STRAIL-Balb/c;
Phalipon A., Costachel C., Thuizat A., Nato F.;
Phalipon A., Nato F.;
Phalipon A., Costachel C., Thuizat A., Nato F.;
Phalipon A., Nat
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Immungolobulin heavy chain variable region (Fragment).
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Phalipon A., Costachel C., Thuizat A., Nato F.;
Phalipon A., Costachel C., Phalipon A., Nato F.;
Phalipon A., Costachel C., Phalipon A., Nato F.;
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25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Immungolobulin heavy chain variable region (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRELIMINARY;
                                                                                                                  AA;
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12863 MW;
                                                                                                                  12833 MW;
           72.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72.2%;
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     Score 70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; E
Sciurognathi; Muridae;
                                                                                                            C85932C0D843778D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5EC11AD02E911952 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   s protective against Shigella
immunodominant serotype-spec:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunodominant
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     DB 2;
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Length 116;
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stepleton M.J., Usdin T.B., Toshiyuki S., Casrinci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., McTley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Krzywinski M.I., Skalska U., Smallus D.E., Schnerch A., Schein J.E.,
A Krzywinski M.I., Skalska U., Smallus D.E., Schnerch A., Schein J.E.,
A Lones S. J. Marra M. A.
                                                                                                                                                      Query Match
Best Local
                                                                                                                     Matches
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099LA6;
01-JUN-2001 (TrEMBLrel. 17,
01-JUN-2001 (TrEMBLrel. 17,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                               SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR007110; Ig-11ke.
InterPro; IPR003597; Ig C1.
InterPro; IPR003006; Ig MHC.
InterPro; IPR003596; Ig v.
Prim; PF07654; C1-set; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN-mix FVB/N; TISSUE-Mammary tumor;
MEDLINE-22388257; PubMed.12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
                                                                                                                                                                                                                                                 Hypothetical 
SEQUENCE 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIR; F33932; F33932.
PIR; S19963; S19963.
HSSP; P01810; 2FBJ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; BC003495; AAH03495.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Strausberg R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-mix FVB/N; TISSUE-Mammary tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   σ
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69
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WVNIETGESVYADDFK 84
                                                       WINTYTGQSTYADDFK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (FEB-2001) to the
                                                                                                                                                                                                                                              ll protein.
484 AA; 52567 MW;
                                                                                                                        Conservative
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                                                                                                                                                   70.1%;
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Last annotation update)
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1; Mismatches
                                                                                                                                                   Score 68; DB 2;
Pred. No. 0.011;
                                                                                                                                                                                                                                                 8EAEA4F9BCF582FA CRC64;
                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        484
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                                                                                                                                                                             Length 484;
                                                                                                                        Indels
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Schein J.E.,
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RESULT 7
RESULT 7
RESULT 05-J
DF 05-J
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M.J., Usdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
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RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robeak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Vilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Touchman J.W., Green E.D., Dickson M.C.,
RA RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                              Q925S1; PRELIMINARY;
Q925S1; O1-DEC-2001 (TrEMBLrel. 19,
01-DEC-2001 (TrEMBLrel. 19,
01-OCT-2003 (TrEMBLrel. 25,
                                                                         Mus musculus (Mouse)
Eukaryota; Metazoa; (
Mammalia; Eutheria; )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Unotabetical rootein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strausberg R.;
Submitted (JUN-2002) to the
EMBL; BC032733; AAH32733.1;
HSSP; P01751; 1A6W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata, Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUL-2004
05-JUL-2004
SEQUENCE FROM N.A
                                             NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical SEQUENCE 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=22388257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Lymph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Generation and initial analysis and mouse cDNA sequences.";
                                                                                                                                                        (Fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                 69 WENTYTGNPTYAQGF 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 WINTYTGQSTYADDF 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    al protein.
125 AA; 13913 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; PubMed=12477932; DOI=10.1073/pnas.242603899;
Feingold E.A., Grouse L.H., Derge J.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sci. U.S.A. 99:16899-16903(2002).
                                                                              Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61.9%;
                                                                                                                                                                              Created)
Last sequence update)
Last annotation update)
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Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 60; DB 2;
Pred. No. 0.061;
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                                                                            Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B76CE434F5A69788 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of more than 15,000
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                                                                                                                                                                                                                                                                                               218 AA
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د.
                                                                                 Euteleostomi;
; Murinae; Mus
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ACCONCINENT OF THE PROPERTY OF
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CRB3_SC
RX MEDLINE=21848401; PubMed=11859360; DOI=10.1038/nature724;
RA WOOD V., Gwilliam R., Rajandream M.A., Lyne M., Lyne R., Stewart A.,
RA Sgouros J., Peat N., Hayles J., Baker S., Basham D., Bowman S.,
RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.N.,
RA Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G.,
RA Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G.,
RA Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G.,
RA Gentles S., Goble A., Hamlin N., Harris D., Hodgson G.,
RA Gollows K., Jones M., Leather S., McDonald S., McLean J.,
RA James K., Jones M., Leather S., McDonald S., McLean J.,
RA James K., O'Neil S., Pearson D., Quail M.A., Rabbinowitsch E.,
RA Mooney P., Moule S., Mungall K., Murphy L., Niblett D., Odell C.,
RA Gollows K., Rutter S., Saunders D., Seeger K., Sharp S.,
RA Rutherford K., Rutter S., Saunders D., Seeger K., Sharp S.,
RA Skelton J., Simmonds M., Squares R., Squares S., Stevens K.,
RA Taylor K., Taylor R.G., Tivey A., Walsh S.V., Warren T., Whitehead S.,
RA Moodward J., Volckaert G., Aert R., Robben J., Grymonprez B.,
RA Weltjens I., Vanstreels E., Rieger M., Schaefer M., Mueller-Abuer S.,
RA Gabel C., Fuchs M., Fritzc C., Holzer E., Moestl D., Hilbert H.,
RA Borzym K., Langer I., Beck A., Lehrach H., Reinhardt R., Pohl T.M.,
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Best Local S
Matches 9
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Q10272;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

MEDLINE=98070339; PubMed=9407031;

Saka Y., Esashi F., Matsusaka T., Mochida S., Yanagida M.;

"Damage and replication checkpoint control in fission yeast is ensured by interactions of Crb2, a protein with BRCT motif, with Cut5 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schizosaccharomyces pombe (Fission yeast).
Sukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
Schizosaccharomycetales; Schizosaccharomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WD-repeat protein crb3.
Name=crb3; ORFNames=SPAC13G7.08c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HSSP; P01665; 1QNZ.
SMART; SM00406; IGv; 1.
PROSITE; P850835; IG LIKE; 1.
NON TER 218 218
SEQÜENCE 218 AA; 23013 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genes
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NCBI_TaxID=4896;
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25-OCT-2004 (Rel. 45, Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=BALB/c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Mechanism of exogenous nucleic acids and the repair of intestinal epithelium after world J. Gastroenterol. 6:709-717(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=BALB/c;
PubMed=11819679;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the
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irradiated mice k
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WINTHSGVPKYAEEFK 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WINTYTGOSTYADDFK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          218 AA; 23013 MW;
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nes related
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elated to repairing
treatment with the
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Pred. No. 0.34;
4; Mismatches
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Q6LZI9;
05-JUL-2004
05-JUL-2004
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                                                                                                                                                  STRAIN=S2 / LL;
Hendrickson E.L., Kaul R., Zhou Y., Bovee D., Conway de Macario E., Dodsworth J., Gillett W., Conway de Macario E., Land M.L., Levy R., Lie Haydock A.K., Kang A., Land M.L., Levy R., Lie Moore B., Forat I., Overbeek R., Palmeiri A., Moore B., Forat I., Call B., Miller B., Barther B., Forat I., Call B., 
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methanogen
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Archaea; Euryarchaeota; Methanococci;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical protein precursor. OrderedLocusNames=MMP0638;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein precursor.
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InterPTO; IPR011046; WD40 like.
Pfam; PF00400; WD40; 4.
PRINTS; PR00320; GPROTEINBRPT.
SMART; SM00320; WD40; 4.
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                                         "Complete genome sequence of the mesophilic
                                                                                  Leigh J.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=39152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Methanococcaceae; Methanococcus
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                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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PS50082; WD_REPEATS_2;
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446 AA;
Methanococcus maripaludis
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Pred. No. 2
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                                                                                                                                                                              Bovee D., Chap.... D.E. Gillett W., Graham D.E. - B. Lie T.J., Major
                                                                                                                       W.B., Larimer F.W.,
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                                                                                                                                Olson M.V.,
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QBA3F1
QB
                                                                                                                                                                                               OrderedLocusNames-plu0899;
Photorhabdus luminescens (subsp. laumondii).
Bacteria; Proteobacteria; Gammaproteobacteria;
                                                                                                                                                                                                                                                                                                                                 Q7N848 PRELIMINARY; PRT; 2009 AA.
Q7N848;
01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Similarities with peptide synthetase like pristinamycin
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STRAIN=VPI-5482 / ATCC 29148;

MEDLINE=22550858; PubMed=12663928; DOI=10.1126/science.1080029;

MEDLINE=20550858; PubMed=12663928; DOI=10.1126/science.1080029;
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Bacteria; Bacteroidetes; Bacteroides (class); Bacteroidales;
Bacteroidaceae; Bacteroides.
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InterPro; IPRO01440; TPR.-like.
InterPro; IPRO08941; TPR-like.
PROSITE; PS50293; TPR_REGION; 1.
Complete proceome; Hypothetical I
SEQUENCE 759 AA; 88928 MW; 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Xu J., Bjursell M.K., Himrod J., Deng S., Chiang H.C., Hooper L.V., Gordon J.I., "A genomic view of the human-Bacteroides t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Science 299:2074-2076(2003
EMBL; AE016938; AAO78109.1
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MEDLINE=22957627;
                                                             SEQUENCE FROM N.A.
                                                                                                                              NCBI_TaxID=141679;
                                                                                                                                                                      Enterobacteriaceae; Photorhabdus.
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ilarity 50.0%;
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PubMed=14528314;
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 51; DB
Pred. No. 13;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Created)
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Pred. No.
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; 18D855215DEEB953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l protein.
52D975F32501538E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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RESULT

Q7YW2

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GG; GO:0003824; F:catalytic activity;
GG; GO:0003824; F:catalytic activity;
GG); GO:0008757; F:s-adenosylmethionin;
GG); GO:0008152; P:metabolism; IEA.
InterPro; IPR000171; AA adenyl_dom.
InterPro; IPR0000871; AMP-bind.
InterPro; IPR0001242; Condensatn.
InterPro; IPR0001242; Condensatn.
InterPro; IPR001601; Methyltransf.
InterPro; IPR001601; Methyltransf.
InterPro; IPR001613; Phappanteth_bind
InterPro; IPR006163; Phappantes S.
InterPro; IPR0001613; Phappantes S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q7YTW2 PRELIMINARY;
Q7YTW2;
01-OCT-2003 (TrEMBLrel. 25, Ct)
01-OCT-2003 (TrEMBLrel. 25, Lt)
01-MAR-2004 (TrEMBLrel. 26, Lt)
Hypothetical protein W08G11.1,
ORFNames=W08G11.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Duchaud E., Rusniok C., Frangeul L., Buchrieser C., Givaudan A., Taourit S., Bocs S., Boursaux-Eude C., Chandler M., Charles J.-F., Dassa E., Derose R., Derzelle S., Freyssinet G., Gaudriault S., Wedigue C., Lanois A., Powell K., Siguier P., Vincent R., Wingate V., Zouine M., Glaser P., Boemare N., Danchin A., Kunst F.; Wingate V., "The genome sequence of the entomopathogenic bacterium Photorhabdus Luminescens.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Complete SEQUENCE
                        Steward C.A.;
Submitted (MAR-1997) to t
EMBL; Z92817; CAB07296.1;
                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN-Bristol N2;
MEDLINE=99069613; PubMed=9851916;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PRO0154; AMPBINDING.
TIGREPAMS; TIGRO1733; AA-aden; 1
PROSITE; PS50075; ACP_DOMAIN; 1.
PROSITE; PS00455; AMP_BINDING; 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE;
PROSITE; PS00012; PHOSPHOPANTETHEINE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nat.
  WormBase; WBGene00012346; W08G11.1
                                                                                                SEQUENCE FROM N.A.
STRAIN=Bristol N2;
                                                                                                                                                                                 Science
                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Nematoda;
Rhabditidae; Peloderinae; Caer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Caenorhabditis elegans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00668; Condensation; ; Pfam; PF00550; PP-binding; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PhotoList; plu0899; -.
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                                                                                                                                                                               investigating biology.";
Science 282:2012-2018(1998).
                                                                                                                                                                                                                                  "Genome sequence of the nematode
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13
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SIMILARITY: Belongs to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1043 WINTYTEOPIALDOLEE 1059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GO:0008757; F:S-adenosylmethionine-dependent methyltransf.
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2009 AA;
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Belongs to the ATP-dependent
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; 229657 MW; 82CD8C883D7813AC
                                                   to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                49.5%;
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Caenorhabditis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
.1.
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                                                   databases
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RESULT 14
Q6N091
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Q8A3Y7
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Best Local S
Matches 8
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Best Local Similarity
Matches 7; Conserv
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InterPro; IPR007110; Ig-11ke.
InterPro; IPR003597; Ig-c1.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003906; Ig_V.
Pfam; PP07654; C1-set; 2.
SMART; SM00409; IG; 4.
SMART; SM00407; IGC1; 3.
SMART; SM00406; IGV; 1.
PROSITE; PS00835; IG_LIKE; 4.
PROSITE; PS00835; IG_MHC; UNKNOWN_2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q6N091;
Q6N091;
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686C02220 (Fragment).
Name=DKFZp686C02220;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WormPep; W08G11.1; CE16558.
InterPro; IPRO04951; DUF268.
Pfam; PF03269; DUF268; 1.
Hypothetical protein.
SEQUENCE 367 AA; 42172 MW
                                                                                    QBA3Y7
QBA3Y7
QBA3Y7
QBA3Y7
01-JUN-2003 (TrEMBLrel. 24, Created)
01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Putative TonB-dependant receptor.
OrderedLocusNames=BT2817;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; BX640625; CAE45779.1; -.
HSSP; P01751; 1A6W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The German Human cDNA Consortium; Wambutt R., Heubner D., Mewes H.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q6N091
          Bacteroides thetaiotaomicron. Bacteria; Bacteroidetes; Bact Bacteroides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical protein.
NON_TER 1
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                               WISAYSGQTYYAQNLQ 102
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nilarity 53.3%;
Conservative :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         500 AA; 54160 MW; 3C423A17D65A41E4 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    48.5%; Score 47; DB 2; Length 500;
43.8%; Pred. No. 38;
tive 5; Mismatches 4; Indels
                                        Bacteroides
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STRAIN-VPI-5482 / ATCC 29148;

STRAIN-VPI-5482 / PUDMed=12663928; DOI=10.1126/science.1080029;

MEDLINE=22550858; PubMed=12663928; DOI=10.1126/science.1080029;

Xu J., Bjursell M.K., Himrod J., Deng S., Carmichael L.K.,

Chiang H.C., Hooper L.V., Gordon J.I.;

"A genomic view of the human-Bacteroides thetaiotaomicron symbiosis.";

Resience 299:2074-2076(2003).

Science 299:2074-2076(2003).

EMBL; AE016937; AA077923.1; -..

GO; GO:0019867; C:outer membrane; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0006872; F:transporter activity; IEA.

GO; GO:0005215; F:transporter activity; IEA.
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SEQUENCE 573 AA; 64637 MW; 414CB2DCF2F14847 CRC64;
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ALIGNMENTS

29-AUG-2000 AAY90823; AAY90823 standard; (first entry) protein; 115 B

520C9 hybridoma VH domain SEQ

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NO:24.

Antigen binding site, immunoglobulin; cancer antigen; immunological; antibody; tumour; human; mucin; cancer; cytostatic; hybridoma; specific binding assay; affinity purification; drug targeting; toxin targeting; imaging; genetic; therapeutic.

Homo sapiens.

US6054561-A.

25-APR-2000

08-FEB-1984; 07-JUN-1995; 84US-00577976. 95US-00483749.

11-JAN-1985; 21-MAR-1986; 08-MAY-1988; 11-AUG-1994; 86US-00842476. 88US-00190778. 94US-00288981

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WPI; 2000-338508/29. N-PSDB; AAA38907.

treating cancer. Monoclonal antibody capable of binding useful for affinity purification, drug to human breast cancer antigen targeting, imaging, and

Disclosure; Fig 12; 57pp; English.

RRSULT 1
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RAYSULT ANYSULT The present invention describes a monoclonal antibody (MAb) (I) that binds to a human breast cancer antigen that is also bound by MAb 454Cl1 and 520C9 (produced hybridoma ATCC HB8484 and HB8696, respectively). Alt described is a hybridoma that produces (I). (I) is useful in specific binding assays, affinity purification, drug or toxin targeting, imaging, and genetic or immunological therapeutics for various cancers. The imaging, Also

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present sequence represents a VH domain derived from a 520C9 hybridoma, which is used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (DUPO ) DU PONT DE NEMOURS & CO E I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY06784 standard; protein; 227 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 115 AA;
                                                                                                       Local Similarity
nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.
hes 6; Conservative
118
                                                  1 RFGFAY 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99
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                                                                                                                                                                                                                        227 AA;
RFGFAY 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGEAY 104
                                                                                                    100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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100.0%; Pr
2+ive 0;
                                                                                                       0;
                                                                                                 Score 34; DB:
Pred. No. 55;
0; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
29;
                                                                                                                                                       2
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                                                                                                                                                       Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                    0
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                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                 0
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RESULT 3
AAWO2280
ALD AAWO
XX
AAWO2280
XX
AAWO2280
XX
AAWO2280
XX
AAWO2280
XX
DT 25-M
DT 25-M
DT 29-O
XX
S20C
XW 520C
XW 520C
XW 520C
XW 520C
XW 50ns
XW in v
XW bind
XX
FH Key
FH Key
FH Pept
FT Pept
XX
PD 09-i
XX
PD 09-i
XX
NO-i
X
밁
                                                                                  Query Match
Best Local Similarity
                                              S
                                                                                                                                                                                                                                                                              Variable heavy (VH) and variable light (VL) genes were cloned from a $20009 hybridoma cDNA library, using probes directed toward the antibody constant and joining regions. A two single chain Fv (sFv) gene was constructed by connecting the VH and VL genes with a Ser rich polypeptide linker. The resulting 52009 two sFv gene, which encodes the present sequence, was inserted into an expression vector, transformed into E. coli, and protein expression induced by the addn. of IPTG to the culture medium. A compsn. comprising a carrier and the 2 sFv protein prod. can be used for in vivo imaging, and drug targetting experiments. The 2 sFv protein prod. is a homodimer, in which both fragments target the same antigen, therefore giving greater binding avidity and longer tissue retention times, compared to individual sFv protein prod. fragments. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                Sequence 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Col 33-36; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Compsns. contg. antigen-targetting antibody fragment constructs comprising dimer of single-chain Fv fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-333194/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oppermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  520C9; anti-c-erbB-2 monoclonal antibody; single chain Fv; sFv; construct; polypeptide linker; C-terminal amino acid sequence; in vivo imaging; drug targetting experiment; homodimer; increased; binding avidity; tissue retention time.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5534254-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR ) CHIRON CORP.
(CREA-) CREATIVE BIOMOLECULES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               520C9 anti-c-erbB-2 two single chain Fv construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
29-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW02280 standard; protein; 243 AA.
99
RFGFAY 104
                                                    RFGFAY
                                                                                                              Conservative
                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ring DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9308-00133804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= linker
                                                                                                                                          100.0%;
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                                                                                                              0;
                                                                                                           Score 34; DB
Pred. No. 58;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Houston LL;
                                                                                                                                   DB 58;
                                                                                                                                                                  2
                                                                                                              0;
                                                                                                                                                                  Length 243;
                                                                                                           0
                                                                                                        Gaps
                                                                                                        0;
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RESULT 4 AAW53170

AXU

AAW53170 standard; protein; 243

A

AAW53170;

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ARBSULT 5
AAWB045
ID AAWB
XX AAWB
AC AAWB
XX AAWB
XX B 520C
XX S 520C
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                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                This represents the protein sequence of a 520C9 sFv' (single chain Fv) construct. This was constructed by connecting the Vh and Vl genes with a DNA sequence encoding a Serine rich linker. 520C9 is a monoclonal antibody useful in targeting c-erbB-2 antigen. This dimeric construct can be used in the methods of invention of imaging a preselected antigen expressed in a mammal. The methods are used in magnetic resonance imaging of c-erbB-2 or related antigens in cancer diagnosis. The biosynthetic constructs have enhanced properties as in vivo targeting agents in comparison with intact monoclonal antibodies or their Fab fragments. The dimeric constructs permit the in vivo targetting of an epitope on an antigen with greater apparent avidity, including greater tumour specificity, tumour localisation and tumour retention properties than that of the Fab fragment having the same CDRS as the construct
                                                                                     520C9 sFv sequence
               Synthetic
                                           antigen-expressing
                                                         520C9 sFv; antigen;
                                                                                                                     28-JAN-1999
                                                                                                                                                                             AAW80424 standard; protein; 243 AA
                                                                                                                                                                                                                                                                                                                                                                         Sequence 243 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Col 33-36; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oppermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-FEB-1992;
07-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigen imaging; single chain Fv; sFv; linker; dimeric; cancer; c-erbB-2; tumour; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         520C9 anti-c-erbB-2 sFv' dimeric construct protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Imaging of antigens in vivo - using dimers of single-chain antibody Fv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHIR ) CHIRON CORP.
(CREA-) CREATIVE BIOMOLECULES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1998-311318/27
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                                                                                                                                                                                                                                                                                                              Similarity 6, Conserv
                                                                                                                                                                                                                                                        REGEAY 104
                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ring DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-00831967.
93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95US-00461838
                                           ; tumour cell; antibody 520C9; targeted delivery;
cell.
                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Huston JS,
                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                Score 34; DE
Pred. No. 58;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Houston LL;
                                                                                                                                                                                                                                                                                                                              58;
                                                                                                                                                                                                                                                                                                                                            <u>ب</u>
                                                                                                                                                                                                                                                                                                                ,
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                                                                                                                                                                                                                                                                                                                                            Length 243;
                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 6
ABW00716
ID ABW0
XX
AC ABW0
DT 15-J
XX
C-20C
XW
C-21L
XW
SFV.
XX
US20
PN US20
PN US20
PF 21-J
XX
XX
O6-F
PR 07-C
PR 06-F
PR 05--;
PR 26-J
XX
XX
XX
XX
XX
HIST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                       06-FEB-1992;
07-OCT-1993;
05-JUN-1995;
26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents an antibody 520C9 sFV. Variable heavy and light sequences of antibody 7520C9 are connected, together with a serine linker, to produce the present single chain Fv gene. The present sequence exemplifies the invention. Dimers of the single chain Fv are used for targeted delivery of drugs or imaging agents (e.g. cytotoxins, prodrugs or 99m-technetium) to antigen-expressing cells, particularly for treatment or diagnosis of tumours (especially of ovary or breast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid encoding single-chain Fv fragment specific for antigens - and having C-terminal tail for crosslinking to form dimer with improved pharmacokinetic properties, used to deliver drugs and imaging agents, especially to tumours.
                            Huston JS,
                                                                                                                                                                                                                                                  21-JUN-2001;
                                                                                                                                                                                                                                                                                                  14-NOV-2002.
                                                                                                                                                                                                                                                                                                                                                US2002168375-A1
                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cell proliferation; cytotoxic agent; drug targetting; single-chain Fv;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    520C9 sFv protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABW00716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABW00716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 243 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Col 35-36; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-023541/02.
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07-OCT-1993;
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                                                                            (CHIR ) CHIRON CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV63399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          н
                            Houston LL,
                                                                                                                       92US-00831967.
93US-00133804.
95US-00462641.
2000US-00558741.
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                                                                                                                                                                                                                                                  2001US-00887853
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93US-00133804.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                         Ring
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                  c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. Such as breast and ovarian tumour cells, which is an approx. 200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). A single chain Fv (sFv) is a covalently linked VH-VL heterodimer which is expressed from a gene fusion including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a binding protein formulation for targetting an epitope on an antigen expressed in mammal. The formulation comprises of dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain. The invention is useful for imaging a preselected antigen in a mammal expressing the antigen. It is also useful for targetting drugs that inhibits cell proliferation and cytocoxic agents that kill cells. The present sequence is 520C9 sFv protein. This sequence is used in the exemplification of the
                                                                                                                                                                                    Claim 4;
                                                                                                                                                                                                                               New single chain Fv imaging or treating
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biosynthetic
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genes connected by a peptide-encoding linker.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 is linked to a human interleukin-2 (IL-2) molecule to make a fusion protein. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cancers particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, renal system, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
                                                              Novel nucleic acid molecule from nonchromoprotein enediyne biosynthetic gene cluster from Micromonospora echinospora useful for conferring calicheamicin resistance on a subject.
                                                                                                                                                                                                                                                                                                                                                   M. echinospora calicheamicin biosynthesis protein CalQ.
                                                                                                                                                                                                                                                          WO200279465-A2
                                                                                                                                                                                                                                                                                Micromonospora echinospora; spp.
                                                                                                                                                                                                                                                                                                                 Calicheamicin biosynthetic gene cluster; aryltetrasaccharide; calicheamicin resistance; nonchromoprotein enediyne; enediyne
                                                                                                                                                                                                                                                                                                                                                                          23-OCT-2003
13-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                           ABU11979;
                                                                                                                                                                                                                                                                                                                                                                                                                                ABU11979 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 7; Fig 9; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel immunoconjugate comprises a humanized
                                                                                                                                                                                                              28-NOV-2001; 2001WO-US044285
                                                                                                                                                                                                                                      10-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Austin
                                                                                                                                                                   (SLOK ) SLOAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                       cheamicin resistance; marrow cell; enzyme.
                                                                                                            2003-092897/08.
)B; ABX56041.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121
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                                          25; Page 107-108;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                267 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kwok CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                        2000US-00724797.
                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                   KETTERING INST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         useful for inhibiting tumor cell growth in vivo anti-p185 antibody linked to an Interleukin-2
                                          179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                  453
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 34;
Pred. No.
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                                                                                                                                                                                                                                                                                                                 enediyne; enediyne
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                                                                                                                                                                                                                                                                                                                            aglycone;
                                                                                                                                                                                                                                                                                                                  resistance;
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encoding

proteins and

The present invention relates to the isolation of the Micromonospora echinospora spp. calcichensis calicheamicin biosynthetic gene cluste

enzymes used in the

biosynthetic production

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Matches

Similarity 6; Conserv

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                                                                                                                                                                                 c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. such as breast and ovarian tumour cells, which is an approx. 200,000 mol. wt. actidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). (Updated on 25-MAR-2003 to correct PN
                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                      Example; Page
                                                                                                                                                                                                                                                                                                                                                                     New single chain Fv imaging or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-FEB-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumour antigen; c-erbB-2; G-FIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence of G-FIT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
07-FEB-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Huston JS,
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6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CREATIVE BIOMOLECULES CETUS ONCOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ46086
                                                                                                            534
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ilarity 100.0%;
Conservative (
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(first en
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                                                                                                                                                                                                                                                                                                                                                                          polypeptide binding to C-erbB-2 tumour antigen - breast or ovarian cancer etc.
  100.0%; 9
100.0%; 1
tive 0;
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Score 34; DB 2; -
Pred. No. 1.2e+02;
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Pred. No.
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                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 453;
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                                                          534;
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RESULT 12
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                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                               The invention comprises the amino acid sequence of proteins from the hyperthermophile Methanopyrus kandleri, the invention also comprises the complete genome from Methanopyrus kandleri. The Methanopyrus kandleri proteins of the invention are useful for enhancing the stability and/or activity of other proteins. The Methanopyrus kandleri genome is useful in a variety of diagnostic and analytical methods. The present amino acid sequence represents a Methanopyrus kandleri protein of the invention.
              Novel human diagnostic protein #5067.
                                         13-FEB-2002 (first entry)
                                                                                                                                                                                                                                                          Sequence 78 AA;
                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acids encoding any of about 1700 Methanopyrus kandleri proteins, and the encoded proteins, useful as a medicaments as diagnostic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-748383/70.
N-PSDB; ADM27081.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-MAR-2002; 2002US-0361742P.
14-MAY-2002; 2002US-0380423P.
16-SEP-2002; 2002US-0410974P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-MAR-2003; 2003WO-US006664.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hyperthermophile; protein stability enhancement; protein activity enhancement.
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                                                                                        ABG05076 standard;
                                                                                                                                                                                                                                                                                                                                                                                     Claim 31; SEQ ID NO 864; 1023pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hyperthermophile Methanopyrus kandleri protein #864.
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                                                                                                                                                                                                      Local Similarity hes 5; Conserv
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                                                                                                                                                         KFGFAY 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pavlov A,
                                                                                        protein; 272 AA.
                                                                                                                                                                                                                     91.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pavlova N,
                                                                                                                                                                                                         1;
                                                                                                                                                                                                                     Score 31;
Pred. No.
                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kozyavkin S;
                                                                                                                                                                                                                     DB 7;
                                                                                                                                                                                                         <u>,,</u>
                                                                                                                                                                                                                                  Length 78;
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RESULT 13
ABG17548
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                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                                                                                                            ABG17548 standard; protein; 300 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 272 AA;
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23-AUG-2000; 2000US-00649167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to isolated polynucleotide (I) and polypeptide (II)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HYSE-)
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DB; AAS69263.
                                                                                                                                                                                                                                                                                                                                    113 RYGFAY 118
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       91.2%;
83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                              Score 31; DB 4;
Pred. No. 2.6e+02;
1; Mismatches 0
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Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder.

Novel human diagnostic protein #17539.

ABG17548;

0;

18-FEB-2002

(first entry

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ABG14667
ID ABG14
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XX Hovel
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XW Humar
XW Homo
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention, note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or (quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The
Homo sapiens
                               Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                             18-FEB-2002
                                                                                                                                                                 ABG14667;
                                                                                                                                                                                                 ABG14667 standard; protein; 309 AA
                                                                                          Novel human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also use in diagnostics as expressed sequence tags for identifying expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polymucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 300 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ftp.wipo.int/pub/published_pct_sequences
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23-AUG-2000;
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)B; AAS81735.
                                                                                                                                                                                                                                                                                           273 RYGFAY 278
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                                                                                        diagnostic protein #14658.
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                                                                                                                           (first entry)
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2000US-00649167.
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83.3%;
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Pred. No. 2.9e+02;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                              Length 300;
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC sequences. (I) is useful as hybridisation probes, polymerase chain (reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymucleotides are also used (C) in diagnostics as expressed sequence tags for identifying expressed (C) is useful in gene therapy techniques to restore normal (C) genes. (I) is useful in gene therapy techniques to restore normal (C) generating antibodies against it, detecting or quantitating a complement. (II) and its binding partners are useful in medical imaging (C) sites expressing (II). (II) and (II) are useful in medical imaging (C) sites expressing (II). (II) are useful for treating disorders (I) polypeptide and polymucleotide sequences have applications in (C) olypeptide and polymucleotide sequences have applications in (C) and to produce other types of data and products dependent on DNA and (C) amino acid sequences of the invention. Note: The sequence data for this (C) patent did not appear in the printed specification, but was obtained in (C) electronic format directly from MIPO at (C) of the wino. int/omb/sublished oct sequences
                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                   biological activity; genetic engineering; hybridisation probe; oligomer;
primer; chromosome mapping; gene mapping; recombinant protein production;
                                                                                                                         Human polypeptide sequence SEQ ID NO:2052
                                                                                                                                                                 12-FEB-2004
                                                                                                                                                                                                                                    ADF59685 standard; protein; 350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 20; SEQ ID NO 45026; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 309 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to isolated polynucleotide (I) and polypeptide (II)
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23-AUG-2000;
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DB; AAS78854.
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2000US-00649167.
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                                                                                                                                                             (first entry)
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83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       Score 31; DB 4;
Pred. No. 3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                        0
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Homo sapiens

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Search completed: April 4, 2005, 15:47:20 Job time: 13.9808 secs
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                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a polypeptide (II) with biological cativity, which is encoded by an isolated polymucleotide sequence (I) control the present invention. Also described: (1) a vector comprising (I); (2) an expression vector comprising (I); (3) a host cell genetically congineered to comprise (I) which is operatively associated with a corregulatory sequence that modulates expression of (I) in the host cell; (C (4) a polypeptide (II) encoded by (I); (5) a composition comprising the polypeptide of (4) and a carrier; (6) an antibody directed against the conjugation of (I) in the conjugation of (I) in a composition comprising the polypeptide of (4); (7) detecting (I) or the polypeptide of (4); (a) compound that binds to the polypeptide of (4); (b) and (c) sample; (b) identifying a compound that binds to the polypeptide of (4); (c) producing the polypeptide of (4); and (I) a collection of (I) conjunclectides (I) can be used as hybridisation probes, (I). The polypurcleotides (I) can be used as hybridisation probes, (I) conjuncted the polypeptide of (4); and (I) are the polypeptide of (4); and (I) a collection of (I) are polypeptides (I) can be used as hybridisation probes, (I) conjuncted the polypeptide of (4); and (I) are the p
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Best Local 9
                                                                                                                                                                                                                        Matches
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N-PSDB; ADF59603.
                                                                                                                                                                                                                                                                                                                                  Sequence 350 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 20; SEQ ID NO 2052; 571pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tang YT,
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                                                                                                            193 RYGFAY 198
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Pred. No. 3.3e+02;
1; Mismatches 0;
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Maximum Match 100%
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Perfect score:
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seq length:
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                                                                                                                                                                                                                                                                                                                                                                                                         Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

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9: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

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10: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-09-66-543-10
US-09-766-543-12
US-09-766-543-12
US-10-424-599-183438
US-10-427-963-106568
US-10-156-761-963
US-10-282-122A-49308
US-10-282-122A-49308
US-10-369-493-2442
US-10-369-493-2442
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           Sequence 6, Appli
Sequence 6, Appli
Sequence 10, Appl
Sequence 12, Appl
Sequence 170, 138,
Sequence 170, 138,
Sequence 106568,
Sequence 9633, Ap
Sequence 9633, Ap
Sequence 204252,
Sequence 204252,
Sequence 204252,
Sequence 204252,
                                                                                                                                                                                                                                                    Description
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US-09-887-853-6
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; Patent No. US20020168375A1
; GENERAL INFORMATION:
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APPLICATION NUMBER: US/09/887,853
FILING DATE: 21-Jun-2001
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/133,804
FILING DATE: CURROWN>
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
                                                                                                                                                                                                              COUNTRY: USA
ZIP: 02109
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department STREET: Exchange Place, 53 State Street CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Huston, James S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 205
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OF INVENTION: Biosynthetic Binding Proteins
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Ring, David B.
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16 US-10-424-599-159963
16 US-10-425-114-63712
15 US-10-425-114-63712
15 US-10-425-114-61010
15 US-10-425-114-51324
10S-09-815-242-11689
10S-10-282-122A-63310
15 US-10-282-122A-63310
15 US-10-282-122A-63310
15 US-10-282-122A-63785
10S-10-282-122A-53118
10S-10-282-122A-51118
10S-10-282-122A-51118
10S-10-282-122A-51118
10S-10-282-122A-51118
10S-10-369-493-12537
10S-09-731-872-278
10S-09-731-872-278
10S-09-731-872-278
10S-09-876-997-282
10S-09-876-997-282
10S-09-876-997-282
10S-10-291-265-238
10S-10-291-265-238
10S-10-291-265-238
10S-10-291-265-238
10S-10-291-265-3360
10S-10-369-493-3466
10S-10-369-493-21603
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US-10-424-599-208245
US-10-424-599-161896
US-10-437-963-132767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 278, App
Sequence 282, App
Sequence 300, App
Sequence 278, App
Sequence 300, App
Sequence 300, App
Sequence 31466, App
Sequence 1127, App
Sequence 1127, App
Sequence 112750,
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Sequence 200245,
Sequence 161896,
Sequence 132767,
Sequence 159963,
Sequence 33660, A
Sequence 63712, A
Sequence 63712, A
Sequence 60100, A
Sequence 60100, A
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TELEFAX: 617-248-7100

FELEPHONE:

617-248-7477

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APPLICANT: AUSTIN, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: King, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PP01679.002
CURRENT APPLICATION NUMBER: US/09/766,543
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                                                           RESULT 3
US-09-766-543-10
Sequence 10, Application US/09766543
Patent No. US20020041665A1
GENERAL IMPORMATION:
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OTHER INFORMATION: 520C9 SFV US-10-683-547-6
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LENGTH: 243
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US/09/558,741
PRIOR FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 07/831,967
PRIOR FILING DATE: 1992-02-06
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: CIBT-P01-130
CURRENT APPLICATION NUMBER: US/10/683,547
CURRENT FILING DATE: 2003-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Huston, J.
APPLICANT: Houston, I
APPLICANT: Ring, D.
APPLICANT: Oppermann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Oppermann, H.
TITLE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.0%; Score 34; local Similarity 100.0%; Pred. No. les 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                            99
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TYPE: amino acid
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Pred. No.
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GENERAL INFORMATION:

APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Kwok, Cheuk S.
APPLICANT: Ring, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PP01679-002
CURRENT APPLICATION NUMBER: US/09/766,543
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 12
                                                                                                                                                                                                                                                                                     RESULT 5
US-10-424-599-183438
; Sequence 183438, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
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Best Local Similarity
""" hes 6; Conserve
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                                                                               APPLICANT: La ROSA Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Chou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 183438
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: H520C9sFv plus
OTHER INFORMATION: linker
                                   TYPE: PRT
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 276
TYPE: PRT
ORGANISM: Artificial Sequence
OTHER INFORMATION: Clone ID: PAT_MRT3847_136658C.1.pep
                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: 520C9 OTHER INFORMATION: humanized single-chain antibody used in the OTHER INFORMATION: IL-2-antibody fusions
                                                                         ENGTH:
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TYPE: PRT
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Pred. No. 51;
0; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
ITITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 179913
LENGTH: 122
TYPE: """
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-437-963-106568
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                                                                                                                                                   NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 106568
LENGTH: 541
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Publication No. US20040123343A1
OTHER INFORMATION: Clone ID: PAT_MRT4530_109C.1.pep-10-437-963-106568
                                                                                                                                                                                                 TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(5321)B CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
                                                                                                                                                                                                                                                                                                                                                               APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
ADDITCANT: Co. V.
                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Glycine max FEATURE:
                                                NAMB/KEY: unsure
LOCATION: (1)..(541)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                   ORGANISM: Oryza sativa FEATURE:
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                                    FEATURE:
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Local Similarity 83.3%;
nes 5; Conservativo
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Li, Ping
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83.3%;
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; Sequence 49308, Application US/10282122A
; Publication No. US20040029129A1
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Best Local Similarity 83.3
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LENGTH: 1015
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APPLICANT: SAKAKI, VOSHIVUKI
APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
CURBERNT NOVEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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APPLICANT: Xu, H.

TITLE OF INVENTION: Identification of Essential Genes in Microorganisms

FILE REFERENCE: ELLTRA.034A

CURRENT APPLICATION NUMBER: US/10/282,122A

CURRENT APPLICATION NUMBER: 00/191,078

PRIOR APPLICATION NUMBER: 60/191,078

PRIOR FILING DATE: 2000-03-21

PRIOR APPLICATION NUMBER: 60/206,848

PRIOR APPLICATION NUMBER: 60/206,848

PRIOR FILING DATE: 2000-05-23

PRIOR FILING DATE: 2000-05-23

PRIOR FILING DATE: 2000-05-26
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APPLICANT:
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APPLICANT:
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APPLICANT:
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CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: JP 2001-204089
PRIOR FILING DATE: 2001-05-30
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PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 15109
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5; Conserva
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Trawick, John
Carr, Grant
Yamamoto, Robert
Foreyth, R.
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HORIKAWA, HIROSHI
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Zyskind, Judith
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lio, Carlos
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Pred. No. 7.5e+02;
1; Mismatches
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Pred. No. 4.1e+02
1; Mismatches
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APPLICATION NUMBER: 60/230,335 FILING DATE: 2000-09-06 APPLICATION NUMBER: 60/230,347

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RESULT 10
US-10-424-599-204252
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PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                          Sequence 204252, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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SEQ ID NO 49308
LENGTH: 1531
TYPE: PRT
ORGANISM: Burkholderia fungorum
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
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NUMBER OF SEQ ID NOS: 78614
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NAME/KEY: MISC FEATURE
LOCATION: (611)...(611)
OTHER INFORMATION: X=any amino acid
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NAME/KEY: MISC_FEATURE
LOCATION: (613)...(613)
OTHER INFORMATION: X=any
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NAME/KEY: MISC_FEATURE
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APPLICATION NUMBER: 60/242,578
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د.
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Pred. No. 1.1e+03;
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_26467C.1.pep
US-10-424-599-204252
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US-09-815-242-10894
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Best Local Similarity
Thehes 5; Conserve
                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Haselbeck
APPLICANT: Ohlsen,
APPLICANT: Zyskind,
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NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 204252
                                                                                                                                                                                                                                      Sequence 10894, Application US/09815242 Patent No. US20020061569A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 2442
LENGTH: 738
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Best Local Similarity 83.3%;
Matches 5; Conservative
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                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052) B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
         APPLICANT: Yamamoto, Robert T.
APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
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CURRENT APPLICATION NUMBER: US/09/815,242
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LOCATION: (1)..(738)
OTHER INFORMATION: unsure at all Xaa locations
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ORGANISM: Glycine max
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                                                                                                                                                                                                                                                                                                                                                   148 RFGFGY 153
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                                                                                                                                                           Haselbeck, Robert
Ohlsen, Kari L.
Zyskind, Judith W.
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                                                                                                     Wall, Daniel
Trawick, John D.
Carr, Grant J.
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83.3%;
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Pred. No. 8.8e+02;
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Pred. No. 1.2e+02;
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848

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US-09-815-242-10894
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SOFTWARE: FRETSEQ for Windows Version
SEQ ID NO 10894
LENGTH: 858
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PRIOR FILING DATE: 2000-09-09
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-72
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                                                                                 PRIOR
PRIOR
PRIOR
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                                                                                                                                                                                                                         PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR FILING DATE: 2000-09-06
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PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 5; Conserv
                                                                                                  APPLICATION NUMBER: 60/253,625
FILING DATE: 2000-11-27
                                      APPLICATION NUMBER: 60/257,931
PILING DATE: 2000-12-22
APPLICATION NUMBER: 60/267,636
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APPLICATION NUMBER: 60/257,931
FILING DATE: 2000-12-22
FILING DATE: 2001-02-09
APPLICATION NUMBER: 60/269,308
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Yamamoto, Robert
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Trawick, John
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Zyskind, Jud
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Pred. No. 1e+03;
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RESULT 15
US-10-424-599-208245
; Sequence 208245, Application US/10424599
; Publication No. US20040031072A1
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TYPE: PRT
; ORGANISM: Enterococcus faecalis
US-10-282-122A-57157
; ORGANISM: Glycine max; FEATURE:
CTHER INFORMATION: Clone ID: PAT_MRT3847_30070C.1.pep US-10-424-599-208245
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                                                                                                                                                               APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
PPLICANT: Zhou Yihua
TITLE OF INVENTION: Soy Nucleic Acid Molecui
TITLE OF INVENTION: Plants and Uses Thereol
FILE REFERENCE: 38-21(5323)B
FULF REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-424-599-220145
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NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3.1
SEQ ID NO 57157
                                                                                  NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 208245
LEUGTH: 65
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 220145
LENGTH: 60
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Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Cao Yongwei
TITLE OF INVENTION: SQu Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
FULL REFERENCE: 38-21(53223)B
FULL REFERENT APPLICATION NUMBER: US/10/424,599
CURRENT FILLING DATE: 2003-04-28
RUMBER OF SEQ ID NOS: 285684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
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APPLICANT: Kovalic David K
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Local Similarity 83.3%;
les 5; Conservative
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Pred. No. 1e+03;
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Thereof for Plant Improvement
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Best Local (
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AC0228 hypothetical protein YPO1870 [imported] - Yersinia pestis (strain CO92) C;Species: Yersinia pestis C;Species: Yersinia pestis C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004 C;Accession: AC0228 C;Accession: AC0228 R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, M.B.; deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, G.; Et il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barrell, Et Nature 413, 523-527, 2001 A,Title: Genome sequence of Yersinia pestis, the causative agent of plague A,Reference number: AB0001; MUID:21470413; PMID:11586360 A; Reference number: AB0001; A; Accession: AC0228 A; Cross-references: UNIPROT: Q8ZF48; GB: AL590842; PIDN: CAC90687.1; PID: g15979892; GSPDB: G A; Residues: 1-48 < KUR> A; Molecule type: DNA A;Status: preliminary Query Match YPO1870 30 l Similarity 5; Conserva 1 RFGFAY 6 RFGFSY 35 Conservative 91.2%; 83.3%; 1; Mismatches Score 31; I Pred. No. 5. ALIGNMENTS DB 2; <u>.</u> Length 48 Indels 0 Gaps 0

C;Species: Trichoderma harzianum C;Date: 10-Sep-1999 #sequence revision 10-Sep-1999 #text_change 09-Jul-2004 C;Accession: S26196; S19787 R;Goldman, G.H.; Demolder, J.; Dewaele, S.; Herrera-Estrella, A.; Geremia, R.A.; van Mont Mol. Gen. Genet. 234, 481-488, 1992 A;Title: Molecular cloning of the imidazoleglycerolphosphate dehydratase gene of Trichode A;Reference number: S26196; MUID:93024323; PMID:1408594 A;Accession: S26196 RESULT 2 \$26196 imidazoleglycerol-phosphate dehydratase (EC 4.2.1.19) - fungus (Trichoderma harzianum)

A;Gene: igh
C;Superfamily: imidazoleglycerol-phosphate dehydratase; imidazoleglycerol-phosphate
C;Keywords: carbon-oxygen lyase; histidine biosynthesis; hydro-lyase
F;58-208/Domain: imidazoleglycerol-phosphate dehydratase homology <IPD> A;Molecule type: mRNA A;Residues: 1-208 <GOL> A;Cross-references: UNIPROT:P34041; EMBL:Z11528; NID:g5176; PIDN:CAA77617.1; PID:g5177 C;Genetice: 91.2%; Similarity 83.3%; 5; Conservation Score 31; Pred. No. 2 DB 1; Length 208; 26;

1; Mismatches

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A;Residues: 1-1008 <SIM>
A;Cross-references: UNIPROT:(OPPEJ2; GB:AE003940; GB:AE003849; NID:g9105966; PIDN:AAF8384
A;Cross-references: UNIPROT:(OPPEJ2; GB:AE003940; GB:AE003849; NID:g9105966; PIDN:AAF8384
A;Experimental source: strain 9a5c
R;Simpson, A.J.G:; Reinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; A
Briones, M.R.S.; Bueno, M.R.P.; Camargo, A.A.; Camargo, L.B.A.; Carraro, D.M.; Carrer, H
as-Neto, E.; Docena, C.; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.
submitted to GenBank, June 2000
A;Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohm
J.D.; Junqueira, M.L.; Kemper, E.L.; Kitajima, J.P.; Krieger, J.B.; Kuramae, B.E.; Laigr
Chado, M.A.; Madeira, A.M.B.N.; Madeira, H.M.F.; Marino, C.L.; Marques, M.V.; Martins, E.A;Authors: Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, E.C.; Miyaki, C.Y.;
F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.R.
Rodrigues, V.; Rosa, A.J. de M.; de Rosa JT., V.B.; de Sa, R.G.; Santelli, R.V.; Sawasak
A;Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva JT., W.A.; da Silva,
A;Reference number: A59328
A;Reference number: A59328
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A; Residues: 1-177 < ALO>
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Gene 204, 201-212, 1997
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C; Accession: T42289
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C;Species: Xylella fastidiosa
C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
C;Accession: G82732
R;Anonymous, The Xylella fastidiosa Consortium of the Organization for Nucleotide Sequent Nature 406, 151-157, 2000
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A;Note: for a complete list of authors see reference number A59328 bel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
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                                                                                                                                                                                                                   88.2%;
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Pred. No. 1.3e+02;
1; Mismatches 0;
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l; Mismatches
                                                                                                                                                                                                                   Score 30; DB 2;
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                                                                                                              C;Species: Bacillus subtilis
C;Date: 05-Dec-1997 #text_change 09-Jul-2004
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: C70025; #Sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: C55220
R;Kunst, F; Ogasawara, N; Moszer, I; Albertini, A.M.; Alloni, G; Azevedo, V
C; Bron, S; Brouillet, S; Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Brouillet, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, S, Bruschi, C.V.; Caldwell, B; Capuano, V.; Carter,
C; Bron, C, C. C.; Capuano, C.V.; C
Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C
iech, J.; Harwood, C.R.; Henaut,
                                                                                                                                                                                                                                                                                                                                                               multidrug-efflux transporter homolog yuxJ - Bacillus N,Alternate names: msyB homolog, pbpD 5'-region
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C70025
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A;Reference number: S725
A;Accession: S73011
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-213 <SMI>
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                                                                                                                                                                                                                                                                       Nature 409, 1007-1011, 2001
A;Authors: Rutter, S.; Seeger, K.; Simon, S.; Simmonds, M.;
A;Title: Massive gene decay in the leprosy bacillus.
A;Reference number: A86909; MUID:21128732; PMID:11234002
A;Accession: G87202
                                                                                                                                                                                                                                                                                                                                                                   R;Cole, S.T.; Eiglmeier, K.; Parkhill, J.; James, K.D.; Thomson, N.R.; Wheeler, R.; Davies, R.M.; Devlin, K.; Duthoy, S.; Feltwell, T.; Fraser, A.; Hamlin, N.; eam, M.A.; Rutherford, K.M.
                                                                                                                                                                                                                                                                                                                                                                                                                               probable membrane protein [imported] - Mycobacterium leprae C;Species: Mycobacterium leprae C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change C;Accession: G87202
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R;Smith, D.R.; Robison, K.
                                                                                                                                                                                                                A; Residues: 1-336 <STO>
                                                                                                                                                                                                                                   A; Molecule type:
                                                                                                                                                                                                                                                    A; Status: preliminary
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C;Superfamily:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      submitted to the EMBL Data Library, November 1993 A; Description: Mycobacterium leprae cosmid L518.
                                                                           Matches
                                                                                                              Query Match
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83.3%;
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Pred. No. 68;
1; Mismatches
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Pred. No.
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D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; stein, G.; Krogh, S.; Kumano, M.; Kurita, K.;

S.; Galizzi, A.; G Hosono, S.; Hullo, Lapidus, A.; Lardi

, A.; Gallero Hullo, M.F.;

, V.; Berterc r, N.M.; Choi Ferrari, E.

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R;Stover, C.K.; Pham, X.Q.; E
adman, S.; Yuan, Y.; Brody, L
; Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
A;Title: Complete genome sequ
A;Reference number: A82950; M
A;Accession: D83625
C;Species: Sc
C;Date: 10-De
C;Accession:
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A;Residues: 101-356 <POP>
A;Cross-references: GB:U11882; NID:g520534; PIDN:AAA64942.1; PID:g520535
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J. Bacteriol. 176, 7197-7205, 1994
A;Title: Cloning, nucleotide sequence, mutagenesis, and A;Reference number: A55220; MUID:95050302; PMID:7961491
A;Accession: C55220
isp4 protein - fission yeast (Schizosaccharomyces pombe) C;Bpecies: Schizosaccharomyces pombe C;Date: 10-Dec-1994 #sequence_revision 21-Jul-1995 #text C;Accession: S45495; S43741
                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:Q916W6; GB:AE004454; GB:AE004091; NID:g9945990; PIDN:AAG0355
A;Experimental source: strain PAO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probable transporter PA0166 [imported] - Pseudomonas aeruginosa
C,Species: Pseudomonas aeruginosa
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Experimental source: strain 168
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Best Local
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50; MUID:20437337; PMID:10984043
                                                                                                                                                                                                                                                                                            88.2%;
83.3%;
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L.L.; (
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Pred. No. 72;
0; Mismatches
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                21-Jul-1995 #text_change 09-Jul-2004
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K.; Lim,
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R;She, Q.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Chan-V Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, P. arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001
A;Description: Sulfolobus solfataricus complete genome.
A;Reference number: A99139
A;Accession: B99358
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B99358
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F90362
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C;Superfamily: Saccharomyces cerevisiae probable membrane protein YJL212c
                                                                                                                                                                                                                                                     C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 09-Jul-2004 C;Accession: B99358
                                                                                                                                                                                                                                                                                                   hypothetical protein SSO9270 [imported] - Sulfolobus solfataricus transposon C_iSpecies: Sulfolobus solfataricus
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A; Residues: 1-98 < KUR>
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Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, P.
arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J.
submitted to GenBank, April 2001
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A; Residues: 1-776 < SAT>
                A; Molecule type: DNA
A; Residues: 1-98 < KUR>
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A; Accession: F90362
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A; Cross-references:
                                                                 A;Status: preliminary
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UNIPROT: Q97X27; GB: AE006641; NID: g13815199; PIDN: AAK42121.1;
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Pred. No. 1.6e+02
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  GSPDB:GI
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Query Match Best Local Similarity

85.3%; 83.3%;

Score Pred.

29; DB No. 32;

<u>ب</u>

Length 98

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Ig heavy chain V region precursor - mouse C;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999 C;Accession: S42467 R;Shiyanov, P.A.; Bespalov, I.A.; Terletskaya, H.N.; Deyev, S.M. submitted to the EMBL Data Library, March 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Parkhill, J.; Wren, B.W.; Mungall, K.; Ketley, J.M.; Churcher, C.; Basham, C.W.; Quail, M.; Rajandream, M.A.; Rutherford, K.M.; VanVliet, A.; Whitehead, Nature 403, 665-668, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             probable integral membrane protein Cj1165c [imported] - Campylobacter jejuni C;Species: Campylobacter jejuni C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 09-Jul-2004 C;Accession: G81321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: mRNA
A;Residues: 1-137 <5HI>
A;Residues: 1-137 <5HI>
A;Cross-references: EMBL:X78107; NID:g460798; PIDN:CAA54997.1; PID:g460799
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;34-119/Domain: immunoglobulin homology <IMM>
R;Bult, C.J.; White, O.; Olsen, G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Blai; Reich, C.I.; Overbeek, R.; Kirkness, E.F.; Weinstock, K.G.; Merrick, J.M.; Glodek, rson, J.D.; Sadow, P.W.; Hanna, M.C.; Cotton, M.D.; Roberts, K.M.; Hurst, M.A.
                                                              hypothetical protein MJ0910 - Methanococcus jannaschii C;Species: Methanococcus jannaschii C;Species: Methanococcus jannaschii C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 09-Jul-2004 C;Accession: F64413
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C; Superfamily:
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A;Molecule type: DNA
A;Residues: 1-164 <PAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hyph;Reference number: A81250; MUID:20150912; PMID:10688204
A;Accession: G81321
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A; Accession: S42467
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A;Experimental source: serotype O2, strain NCTC 11168
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100.0%; Pred. No. 45;
tive 0; Mismatches
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Science 273, 1058-1073, 1996
A,Anthors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.; Woese, C. A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschii. A;Reference number: A64300; MUID:96337999; PMID:8688087
A,Accession: F64413
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E90356
                                                                                                                                                                                                                                                                                                                                                               R;She, Q.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Cha: Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001
A;Description: Sulfolobus solfataricus complete genome.
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Search completed: April Job time: 3.16613 secs
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A; Accession: E90356
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5; Conservative
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57;
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re greater than or equal to the score of the result being printed,
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Q978d3 wolbachia
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Q92h44 wolbachia
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Q915y7 wolbachia
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Q841p2 wolbachia
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Q947p3 wolbachia
Q95p48 yersinia p
Q86p48 yersinia p
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Q66p59 yersinia p
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Q915Y6;
01-CCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
0uter surface protein (Fragment).

Name=wsp;
Wolbachia endosymbiont of Trichogramma embryophagum.
Bacteria; Proteobacteria; Alphaproteobacteria; Ricketts
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STRAIN=Uro3;

STRAIN=20496957; PubMed=11040288;

MEDLINE=20496957; PubMed=7., Lassabliere F., Cha
Pintureau B., Chaudier S., Lassabliere F., Cha
Pintureau B., Chaudier S., Lassabliere F., Cha
"Addition of wsp sequences to the Wolbachia ph
"Addition of wsp sequences to the Wolbachia ph
stability of the classification.";

J. Mol. Evol. 51:374-377(2000).

EMBL; AF245165; AAF71319.1;

InterPro; IPR002566; Surface Ag_msp4.
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                                                                                                                        Wolbachia sp. wBtab6.
Bacteria; Proteobacteria; Alp
Bacteria; Proteobacteria; Alp
Rickettsiaceae; Wolbachieae;
RCBI_TaxID=133462;
                                                                                                                                                                                                                                                        Name=wsp;
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   MEDLINE=22867875; PubMed=14506854;
                                    SEQUENCE FROM N.A. STRAIN=wBtab6;
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Q8PEBG9
Q8PEBG9
Q8PEBG9
Q8PEBG9
Q8PEBG9
Q8PEBG9
Q8PEBG9
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Q8WFBBC38
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Q6C7r4 yarrowia li
Q8egg7 shewanalla
Q87ei9 xylella fas
Q9pej2 xylella fas
Q8pej5 xylella fas
Q8per6 xanthomonas
Q8per6 xanthomonas
Q8per6 satreptomyce
Q821b5 streptomyce
Q70k09 bactilus am
Q48449 bactilus am
Q4849 bacteriopha
Q8wy49 homo sapien
Q49930 pseudomonas
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Q93IJ3;
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation updat
Outer surface protein (Fragment).
                                                                Wolbachia sp. wBtab3.
Bacteria; Proteobacteria; Alphaproteobacteria;
Rickettsiaceae; Wolbachieae; Wolbachia.
NCBI_TaxID=133459;
STRAIN=wBtab3;
                        SEQUENCE FROM N.A.
                                                                                                                                                                Name=wsp;
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EMBL; AJ291375; CAC44875.1; -.
InterPro; IPR002566; Surface Ag_msp4.
Pfam; PF01617; Surface Ag_2; 2.
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Nirgianaki A., Banks G.K., Frohlich D., Veneti
Miller T.A., Bedford I.D., arkham P.G., Savak
"Wolbachia infections of the whitefly Bemisia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q93IJ2; PRELIMINARY; PRT; 185 AA.
Q93IJ2;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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EMBL; AJ291377; CAC44877.1; -.
InterPro; IPR002566; Surface_Ag_msp4.
Pfam; PF01617; Surface_Ag_2; 2.
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Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
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emisia tabaci.";
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Bourtzis K.;
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Best Local :
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Q9RBD3;
Q1-WAY-2000 (TrEMBLrel. 13, Created)
O1-WAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-UNN-2002 (TrEMBLrel. 21, Last annotation update)
O1-Unrenorface protein (Fragment).
                                                                Wolbachia sp. wDeiSW.
Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
Rickettsiaceae; Wolbachieae; Wolbachia.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence up
01-UN-2002 (TrEMBLrel. 21, Last annotation
Outer surface protein (Fragment).
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Mirgianaki A., Banks G.K., Frohlich D., Veneti D.R., Braig H.:
Miller T.A., Bedford I.D., arkham P.G., Savakis C., Bourtzis:
"Wolbachia infections of the whitefly Bemisia tabaci.";
Curr. Microbiol. 47:93-101(2003).
EMBL; AJ291374; CAC44874 1; -
InterPro; IPR002566; Surface_Ag_msp4.
Pfam; PF01617; Surface_Ag_2; 2.
  SEQUENCE FROM N.A.
                                       NCBI_TaxID=77831;
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EMBL; AF071926; AAC77406.1; -.
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Matches 6
Q9ZH45 PRELIMINARY; PRT; 185 AA.

Q9ZH45;

Q9ZH45;

T01-MAY-1999 (TYEMBLrel. 10, Created)

T01-MAY-1999 (TYEMBLrel. 10, Last sequence update)

T01-UN-2002 (TYEMBLrel. 21, Last annotation update)

Quter surface protein (Fragment).

Name=wSp;

Wolbachia sp. wKayB.

Bactearia; Proteobacteria; Alphaproteobacteria; Ricke Rickettsiaceae; Wolbachia.

C Rickettsiaceae; Wolbachia.

X NCBI TaxID=77830;

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STRAIN=wKmyLC;
MEDLINE=99398205; PubMed=10469257;
van Meer M.M.M., Witteveldt J., Stouthamer R.;
"Phylogeny of the arthropod endosymbiont Wolbachia based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wolbachia sp. wKayLC.
Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales; Rickettsiacea; Wolbachieae; Wolbachiea
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01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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InterPro; IPR002566; Surface_Ag_msp4
Pfam; PF01617; Surface_Ag_2; 2.
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Pfam; PF01617; Surface_Ag_2; 2.
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EMBL; AF071925; AAC77405.1; -.
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Pred. No. 34;
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Pred. No. 34;
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RESULT

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ID ZH46

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AC Q
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Matches 6
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Q9L5Y4;
Q1-CT-2000 (TrEMBLrel. 15, Created)
Q1-CT-2000 (TrEMBLrel. 15, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Quter surface protein (Fragment).
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Q9ZH46;
Q1-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Outer surface protein (Fragment).
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Rickettsiaceae; Wolbachieae; NCBI_TaxID=125597;
                                                     Wolbachia endosymbiont of Trichogramma evanescens
Bacteria; Proteobacteria; Alphaproteobacteria; Ri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=99398205; PubMed=10469257; van Meer M.M.M., Witteveldt J., St
"Phylogeny of the arthropod endosy
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EMBL; AF071924; AAC77404.1; -
InterPro; IPR002566; Surface Ag_msp4.
Pfam; PF01617; Surface_Ag_2; 2.
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InterPro; IPR002566; Surface_
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Pred. No. 34;
0; Mismatches
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Pred. No. 34;
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Wolbachia based
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                                                          Rickettsiales;
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RESULT 12
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Best Local 9
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Q9L5Y7;
Q1-5CT-2000 (T1
01-0CT-2000 (T3
01-JUN-2002 (T1
Outer surface F
  Wolbachia endosy
Bacteria; Proteo
Rickettsiaceae;
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SEQUENCE FR
STRAIN=M36;
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EMBL, AF245167, AAF71321.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF
NON_TER
NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wolbachia endosymbiont of Trichogramma oleae.
Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
Rickettsiaceae; Wolbachieae; Wolbachia.
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01-OCT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 15,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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                                                                        Name=wsp;
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Pfam; PF01617; Surface_Ag_2; 2.
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InterPro; IPR002566; Surface_Ag_msp4
Pfam; PF01617; Surface_Ag_2; 2.
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    endosymbiont of Trichogramma cordubensis.
Proteobacteria; Alphaproteobacteria; Rickettsiales;
aceae; Wolbachieae; Wolbachia.
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186 AA;
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ace protein (Fragment).
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ilarity 100.0%;
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20144 MW;
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Pred. No. 35;
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Pred. No.
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a phylogenetic tree and
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O915Y8
O1-OCT-2000 (TrEMBLrel. 15, Created)
O1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
O1-OCT-2000 (TrEMBLrel. 21, Last annotation update)
O1-TUN-2002 (TrEMBLrel. 21, Last annotation update)
On-ter surface protein (Fragment).
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"Addition of wsp sequences to the Wolbachia phylogenetic tree and
stability of the classification.";
J. Mol. Evol. 51:374-377(2000).
EMBL; AF245164; AAF71318.1.;
InterPro; IPR002566; Surface Ag_msp4.
Pfam; PF01617; Surface Ag_msp4.
NON TER 186 186
                                                      Q9L5Y9 PRELIMINARY;
Q9L5Y9; PRELIMINARY;
Q1-CT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 21,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                    MEDLINE=20496957; PubMed=11040288;
Pintureau B., Chaudier S., Lassabliere F., Charles H., Grenier S.;
Pintureau B., Chaudier S., Lassabliere F., Charles H., Grenier S.;
Paddition of wps sequences to the Wolbachia phylogenetic tree and stability of the classification.";
J. WOL. Evol. 51:374-377(2000).
EMBL; AF245163; AAF71317.1; -.
InterPro; IPR002566; Surface_Ag_msp4.
Pfam; PF01617; Surface_Ag_2; 2.
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Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
Rickettsiaceae; Wolbachieae; Wolbachia.
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                    Wolbachia
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         Bacteria;
                                           Outer surface protein (Fragment).
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         endosymbiont of Trichogramma semblidi
Proteobacteria; Alphaproteobacteria;
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186 AA;
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tive 0; Mismatches
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                                                                                                           186
semblidis.
~~*eria; Rickettsiales;
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RESULT 15
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AC Q6GV1
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MEDLINE=20496957; PubMed=11040288;

MIDILINE=20496957; PubMed=11040288;

Pintureau B., Chaudier S., Lassabliere F., Charles H., Grenier S.;

Pintureau B., Chaudier S., Lassabliere F., Charles H., Grenier S.;

"Addition of wsp sequences to the Wolbachia phylogenetic tree and stability of the classification.";

J. Mol. Evol. 51:374-377(2000).

EMBL, APF45162; AAF71316.1;

InterPro; IPR002566; Surface Ag_msp4.

Pfam; PF01617; Surface Ag_msp4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BEQUENCE FROM N.A.

BERMANG C.M., Cong B., Dai Q.H., Fu H.B.;

Wang C.M., Cong B., Dai Q.H., Fu H.B.;

Wang C.M., Cong B., Dai Q.H., Fu H.B.;

Submitted (MAY-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AY634680; AAT48492.1; -.

InterPro; IPR002566; Surface Ag_msp4.

Pfam; PF01617; Surface Ag_2; 1.
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05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
outer surface protein (Fragment).
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SEQUENCE
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SEQUENCE
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Bacteria; Proteobacteria; Alphaproteobacteria; Rickettsiales;
Rickettsiaceae; Wolbachieae; Wolbachia.
NCBI_TaxID=164403;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rickettsiaceae, Wolbachieae, Wolbachia.
NCBI_TaxID=125592;
[1]
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186 AA;
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20158 MW;
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Copyright (c) 1993 - 2005
geneseqp1980s:*
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Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Regult No.	Score	ימא	Length	2 B	ID	Description
	1258 1258	100.0	243 243	NN	AAW02280 AAW53170	Aaw02280 Aaw53170
ω	1258	100.0	243	N	AAW80424	Aaw80424
4	1258		243	7	ABW00716	Abw00716
υ.	1258		246	2	AAR39569	Aar39569
6	1206	95.9	534	N	AAR39571	Aar39571
7	1055.5	83.9	409	4	AAU04945	Aau()4945
8	976	77.6	499	s	AAU72872	Aau72872
9	950.5	75.6	267	4.	AAU04944	Aau04944
10	914	72.7	250	N	AAW02278	Aaw()2278
11	914	•	250	N	AAW29261	Aaw29261
12	914	•	250	ผ	AAW22400	Aaw22400
13	914	•	250	N	AAW47012	Aaw47012
14	914	72.7	250	N	AAW53168	Aaw53168
15	914	72.7	250	N	AAW80422	Aaw80422
16	914	72.7	250	7	ABW00714	Abw00714
17	879	69.9	667	N	AAR39573	Aar39573
18	857.5		237	w	AAY44595	Aay44595
19	•	68.2	243	w	AAY44596	Aay14596
20	857		259	7	ADG32322	Adg32322
21	857		329	7	ADG32359	Adg32359
22	842		259	7	ADG32324	Adg32324
23	842		329	7	ADG32361	Adg:32361
24	841.5		240	œ	ADN16154	Adn:16154
25	840.5		242	^	2200023	Aab20433

815.5 64.8 320 7	815.5 64.8 251 7	816 64.9 259 3	65.0 252 5	818.5 65.1 240 2	65.1 240 2	820.5 65.2 637 2	823.5 65.5 1020 2	823.5 65.5 899 2	823.5 65.5 895 2	823.5 65.5 892 2	823.5 65.5 892 2	823.5 65.5 892 2	823.5 65.5 711 2	823.5 65.5 699 2	823.5 65.5 651 2	823.5 65.5 617 2	823.5 65.5 615 2	023.3 03.0 03.0 4
ADG32358	ADG32321	AAB09779	AAU72863	AAW05134	AAR85494	AAR26982	AAW05141	AAW05144	AAW05142	AAW05139	AAW05143	AAW05140	AAR26980	AAW05138	AAW05136	AAR95057	AAR95056	CCOCKNAM
Adg32358	Adg32321	Aab09779	Aau72863	Aaw05134	Aar85494	Aar26982	Aaw05141	Aaw05144	Aaw05142	Aaw05139	Aaw05143	Aaw05140	Aar26980	Aaw05138	Aaw05136	Aar95057	Aar95056	AGLACOCA
Precursor	Mouse scF	Antiviral	P4-14 sin	Single ch	ErbB-2-ap	(FRP5)-ET	scFv2 (FRP	BCFV2 (FRP	scrv2 (FRP	scFv2 (FRP	scFv2 (FRP	BCFv2 (225	Fv (FRP5) -	SCFV (FRP5	BCFV (FRP5	GAL4-DT-8	GAL4-DT-8	OCE A VENER

ALIGNMENTS

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RESULT 1
AAW0220
ID AAW02
XX AAW0
XX AAW0
XX AAW0
XX POT 25-M
DT 29-O
XX S20C
XX S20C
XX S20C
XX CONB

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     520C9 anti-c-erbB-2 two single chain Fv construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
29-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW02280;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
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eide 334254-A. 7UL-1996. CTT-1993; 9	Key Location/Qual	
118. 133 /label= linker 4-A. 996. 993, 93US-00133804. 992, 92US-00831967.	Location/Qualifiers	

Oppermann H, Ring DB, Huston JS, Houston LL;

N-PSDB; 1996-333194/33. DB; AAT36880.

Compans. contg. antigen-targetting antibody fragment constructs comprising dimer of single-chain Fv fragments.

Example 1; Col 33-36; 30pp; English.

Variable heavy (VH) and variable light (VL) genes were cloned from a \$2000 hybridoma cDNA library, using probes directed toward the antibody constant and joining regions. A two single chain Fv (8FV) gene was constructed by connecting the VH and VL genes with a Ser rich polypeptide linker. The resulting 52009 two sFv gene, which encodes the present

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RESULT 2
AAW53170
ID AAW5
XX AAW5
XX AAW5
XX AAW5
XX AAT1
XX AAT1
XX AAT1
XX AAT1
XX AAT1
XX US57
XX US57
XX US57
XX O6-F
PR 07-C
XX OCHI
PA (CHI
PA (
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Best Local
Example 1; Col 33-36; 30pp; English
                                                       Imaging of antigens in fragments.
                                                                                                                                    WPI; 1998-311318/27.
N-PSDB; AAV21798.
                                                                                                                                                                                                              Oppermann H,
                                                                                                                                                                                                                                                                     (CHIR )
                                                                                                                                                                                                                                                                                                                                                   06-FEB-1992;
07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-MAY-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antigen imaging; single chain Fv; tumour; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               520C9 anti-c-erbB-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence, was inserted into an expression vector, transformed into E. coli, and protein expression induced by the addn. of IPTG to the culture medium. A compsn. comprising a carrier and the 2 SFV protein prod. can be used for in vivo imaging, and drug targetting experiments. The 2 SFV protein prod. is a homodimer, in which both fragments target the same antigen, therefore giving greater binding avidity and longer tissue retention times, compared to individual SFV protein prod. fragments. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW53170;
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                                                                                                                                                                                                                                                             CHIRON CORP.
CREATIVE BIOMOLECULES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44
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                                                                                                                                                                                                            Ring DB,
                                                                                                                                                                                                                                                                                                                                             92US-00831967.
93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                                          95US-00461838.
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                                                                             vivo - using
                                                                                                                                                                                                              Huston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sFv; linker; dimeric; cancer; c-erbB-2;
                                                                               dimers
                                                                                                                                                                                                              Houston
                                                                          of single-chain antibody
                                                                                                                                                                                                            E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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RESULT 3
AAW80424
ID AAW8
XX AAW8
AC AAW8
AC AAW8
AC S20C
XX 520C
XX 520C
XX 520C
XX 9xnt
XX 9xnt
XX 0559
PN US58
XX US58
PN US58
YN US58
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Best Local Similarity
Matches 243; Conserv
                                                                                                                                                                         06-FEB-1992;
07-OCT-1993;
                    WPI; 1999-023541/02
                                                                                                                                                                                                                                         05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             520C9 sFv sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This represents the protein sequence of a 520C9 sFv' (single chain Fv) construct. This was constructed by connecting the Vh and Vl genes with a DNA sequence encoding a Serine rich linker. 520C9 is a monoclonal antibody useful in targeting c-erbB-2 antigen. This dimeric construct can be used in the methods of invention of imaging a preselected antigen expressed in a mammal. The methods are used in magnetic resonance imaging of c-erbB-2 or related antigens in cancer diagnosis. The biosynthetic constructs have enhanced properties as in vivo targetting agents in comparison with intact monoclonal antibodies or their Fab fragments. The dimeric constructs permit the in vivo targetting of an epitope on an antigen with greater apparent avidity, including greater tumour specificity, tumour localisation and tumour retention properties than
                                                               Oppermann H,
                                                                                                                                                                                                                                                                                    17-NOV-1998
                                                                                                                                                                                                                                                                                                                             US5837846-A
                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                               antigen-expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                    520C9 sFv;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW80424 standard; protein;
                                                                                                                                  (CREA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW80424;
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                                                                                                         CREATIVE BIOMOLECULES CHIRON CORP.
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93US-00133804
                                                                                                                                                                                                                                         95US-00461386
                                                                                                                                                                                                                                                                                                                                                                                                                  ; tumour cell; antibody 520C9; targeted delivery; cell.
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                                                               Huston
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07-OCT-1993;
05-JUN-1995;
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                                                                                                                                                                                                                                                                   21-JUN-2001; 2001US-00887853.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cell proliferation; cytotoxic agent; drug targetting; single-chain Fv;
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Matches 243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel binding protein formulation for targeting epitope on antigen in mammal, comprising dimeric biosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain.
                                                                                                                                                                                                                                                                                                                                 Single chain biosynthetic
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07-FEB-1994
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                                                                                                                                                                                                                                                                                                                              Fv polypeptide; VH-VL heterodimer; immunoglobulin; single polypeptide chain binding site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                           92US-00831967
                                                                                                                                          93WO-US001055
                                                                                                                                                                                                                                                                                                                                                                                                        957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein;
                                                                                                                                                                                                                                                                                                                                                                                                      protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 1258;
Pred. No. 8.8
0; Mismatches
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hes 0;
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RESULT 6
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Best Local S
Matches 243
                                     06-FEB-1992;
                                                                             05-FEB-1993;
                                                                                                                                                                                                          Synthetic
                                                                                                                        19-AUG-1993
                                                                                                                                                             WO9316185-A2
                                                                                                                                                                                                                                                Tumour antigen; c-erbB-2; G-FIT.
                                                                                                                                                                                                                                                                                          Sequence of
                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
07-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. such as breast and ovarian tumour cells, which is an approx.
200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). A single chain Fv (sFv) is a coval-lently linked VH-VL heterodimer which is expressed from a gene fusion including VH- and VL- encoding genes connected by a peptide-encoding linker. Such linker sequences are set forth in AA residues 116-135 in AAR39569, which for the 520C9 monoclonal antibody, a single chain polypeptide can be produced having a binding affinity for a c-erbB-2 related antigen. 'X' in AAR39569 refers to the location of a stop codon in AAQ46084. (Updated on 55-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                  AAR39571;
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR39571 standard; protein; 534 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 60-61; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New single chain Fv imaging or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Huston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          181 IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLEIK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            243;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAD 243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SGSSSSGSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCBASQDIGNSLTWLQQEPDGTTKRL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS
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                                                                                                                                                                                                                                                                                          G-FIT
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(first en
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                                     92US-00831967
                                                                             93WO-US001055
                                                                                                                                                                                                                                                                                                                                entry)
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Pred. No. 8.9e-88;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB,
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RESULT 7
AAU04945
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AC AAU0
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DT 06-A
DT 24-C
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Huma
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Huma
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Homc
OS Mus
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Best Local S
Matches 233
                                                                                                                     Mus sp.
Synthetic.
Chimeric.
                             Protein
                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                            Humanised;
                                                       Protein
                                                                                Peptide
                                                                                                                                                                                            cancer; tumour;
                                                                                                                                                                                                                                  Humanised anti-p185 antibody/IL-2 fusion protein
                                                                                                                                                                                                                                                               06-AUG-2003
24-OCT-2001
                                                                                                                                                                                                                                                                                                          AAU04945;
                                                                                                                                                                                                                                                                                                                                   AAU04945 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    c-erDB-2 refers to a protein antigen expressed on the surface of tumour cells, such as breast and ovarian tumour cells, which is an approx. 200,000 mol. wt. acidic glycoprotein having an isolelectric pt. of about 5.3 (see AAQ46083, AAR39568). (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 65-68; 87pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Huston JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CREA-) CREATIVE BIOMOLECULES (CETU) CETUS ONCOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               470
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  al Similarity
233; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         534 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                  IKRAD 243
                                                                                                                                                                                           antibody; p185; 520C9H; interleukin-2; IL-2;
mour; adenocarcinoma; fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Houston
                                                                                                                                                                                                                                                              (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         534
                       23. .409
/label= Mature_fusion_protein
Location/Qualifiers
                                                                /label=_Signal_peptide
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95.1%;
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Pred. No. 2e-83;
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
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                                                                                                                                                                                                      immunoconjugate;
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Best Local S
Matches 199
                                                                                                                                   The sequence represents a humanised anti-p185 (520C9) single chain antibody/human interleukin-2 (IL-2) fusion protein. The fusion protein (or immunoconjugate) is used to inhibit the growth of tumours or cancer particularly those characterised by overexpression of p185 e.g. human adenocarcinomas and malignant and/or benign tumours of the breast, renselystem, salivary gland, gastrointestinal tract or gastric tumours. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                  Sequence
                                                                                                                                                                                                                        Claim 13; Fig 11;
                                                                                                                                                                                                                                                       Novel immunoconjugate comprises a humanized
                                                                                                                                                                                                                                                                                                                       Austin R,
                                                                                                                                                                                                                                                                                                                                                       (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                     20-JAN-2000; 2000US-0177258P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein
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                                                                          al Similarity
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            61
                                  23
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HAMILTON CIVIC HOSPITALS
HAMILTON REGIONAL CANCER
                                                                                                                   409
ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS
                                           EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWM;WINTYTGQSTY
                                EIQLVQSGPEVKKPGASVKISCKASGYTFANYGMNWMKQAPGKGLEWMGWINTYTGQSTY
                                                                                                                                                                                                                                                                                                                       Kwok CS,
                                                                          Conservative
                                                                                                                  $
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label= CDR
/note= "Complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label- CDR
/note- "Complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "Encode by ACA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note= "Links the antibody to the IL-2 molecule"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           label-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           label= Synthetic/
note= "Links the"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             notes "Complementarity
                                                                                                                                                                                                                       74pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "Encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       abel= CDR
                                                                                                                                                                                                                                                                                                                     Ring
                                                                                   83.9%;
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                                                                                                                                                                                                                                                       useful for inhibiting tumor cell growth in vivo anti-p185 antibody linked to an Interleukin-2
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"Complementarity
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                                                                        Score 1055.5;
Pred. No. 4.3e
L9; Mismatches
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the heavy chain to the light chain"
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1.3e-72;
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           The invention relates to a multifunctional polypeptide comprising a CC domain with a binding site that specifically recognises an extracellular CC group of the NKG2D receptor complex and a second domain which functions CC as a receptor or ligand. The polypeptide and its associated CC composition for the treatment of cancer, infections and/or autoimmune CC composition for the treatment of cancer, infections and/or autoimmune CC compositions. The cancer may be a tumour of the head and neck stomach, CC protozon colon, liver, intrahepatic bile ducts, pancreas, lung, CC larynx, breast, ovary, uterus, cervix, prostate, kidney, testis, thyroid, CC larynx, breast, ovary, uterus, revix, prostate, kidney, testis, thyroid, CC protozon or helminths. The autoimmune diseases, bacteria, fungi, CC protozon or helminths. The autoimmune diseases include multiple CC sciencis, Garave's disease, ankylosing spondylitis, acute anterior CC uveitis, Goodpaseure's syndrome, myasthenia gravis, insulin-dependent CC diabetes mellitus, rheumatoid arthritis, pemphigus vulgaris and autoimmune hepatitis. Sequences ANU728075 represent the NKG2D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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Mayer
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                                                                                                                                                                                                                                                                                                                         Example 5; Fig 16; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                            Multifunctional polypeptides comprising binding sites that specifically recognize extracellular groups of the NKG2D receptor complex and domains which function as receptors or ligands, useful for treating cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo
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Hofmeister R
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RESULT 9
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ID ANU04944
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AX
AAU0
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DT 06-A
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Matches 199; Conserv
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                                                                                      Peptide
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Synthetic.
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24-OCT-2001
                                                                                                                                                                                                                               Chimeric.
                                                                                                                                                                                                                                                                                    cancer;
                                                                                                                                                                                                                                                                                               Humanised;
                                                                                                                                                                                                                                                                                                                    Humanised anti-p185 single chain antibody, 520C9H
                                                                                                                                                                                                                                                                                                                                                                                AAU04944;
                                                                                                                                                                                                                                                                                                                                                                                                    AAU04944 standard; protein;
                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EVQLLESGGGLVQPGGSLKLSCAASGFDFSRYWMSWVRQAPGKGLEWIGEINPDSSTINY 307
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                                                                                                                                                                                                                                                                                  antibody; p185; 520C9H; interleukin-2; IL-2; immunoconjugate;
mour; adenocarcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241
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                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
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/label= CI
                   /note= "Complementarity determining region"
202. .208
                                                                                                                                                         /label= CDR
/note= "Complementarity
72...88
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                                                                                            /label= CDR
/note= "Complementarity determining region"
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/note= "Complementarity
                                          /label= CDR
                                                               note= "Links
                                                          38. 152
label= Synthetic_peptide_linker
label="'**" the heavy chain to the
                                                     .186
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RESULT 10
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AC AAW02
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DT 25-MA

AAW02278

standard;

protein; 250

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AAW02278; 25-MAR-2003

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Best Local
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                                                                                                                                                                                                                                                                                                Sequence 267
                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 7; Fig 9; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel immunoconjugate comprises a humanized
                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptide.
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N-PSDB; AAS09507.
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(HAMI-)
(HAMI-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JUL-2001.
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                      241 RAD 243
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                                                                                                                                                                                                 23
                                                                                                                                                                                                                                                            n 75.6%;
Similarity 71.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HAMILTON CIVIC HOSPITALS RES DEV INC. HAMILTON REGIONAL CANCER CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHIRON CORP.
                                                                                                                                                           ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQCTLVSVSASISSS 120
                                                                                                                                                                                             GSE 262
                                                          IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLEIK 240
                                                                                                           SGSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDCTIKRL
                                               IYATSSLDSGVPSRFSGSGSGTDFTFTISSLQPEDIATYYCLQYAIFPYTFGQGTRLEIK 259
                                                                                              GGSGGGGSGGGSDIQMTQSPSSLSASVGDRVTITCRASQDIGNSLTWYQQKPGKTPKLL
                                                                                                                                              ADDFKERVIMITDISTSTAYMDLRSLRSDDTAVYYCARRFGFAYWGQGTLVIVS---SGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kwok CS,
                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                ΑĄ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Cc
241. .249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label= CDR
/note= "Complementarity determining region"
260, ..267
/label= Glu_Glu_epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for inhibiting tumor cell growth in vivo anti-p185 antibody linked to an Interleukin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Complementarity determining region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB;
                                                                                                                                                                                                                                                32;
                                                                                                                                                                                                                                               Score 950.5; DB 4
Pred. No. 2.5e-64;
32; Mismatches 34
                                                                                                                                                                                                                                                                      DB 4;
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                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                        267;
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29-OCT-1996

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                                                                                                                                                                                                                                                                                     parental 741F8 hybridoma line. A two single chain Fv (8Fv) gene was constructed by connecting the VH and VL genes with a DNA sequence endcoding a polypeptide linker. A synthetic DNA duplex encoding the Cterminal maino acid sequence, (G1y)4-Cys was inserted, and the resulting 741F8 anti-c-erbB-2 two sFv inserted into an expression vector. The resulting gene, which encodes the present sequence, was transformed into E. coll, and protein expression induced by the addn. of IPTG to the culture medium. A compsn. comprising a carrier and the 2 sFv protein prod. can be used for in vivo imaging, and drug targetting experiments. The 2 sFv protein prod. Is a homodimer, in which both fragments target the same antigen, therefore giving greater binding avidity and longer tissue retention times, compared to individual sFv protein prod. fragments. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    741F8; anti-c-erbB-2; monoclonal antibody; single chain Fv; sFv; construct; polypeptide linker; C-terminal amino acid sequence; in vivo imaging; drug targetting experiment; homodimer; increased, binding avidity; tissue retention time.
                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The variable heavy (VH) and variable light (VL) cerbB-2 monoclonal antibody (MAb), were isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Compans. contg. antigen-targetting antibody fragment constructs comprising dimer of single-chain Fv fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oppermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1996-333194/33.
                                117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25; Col 27-28; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     anti-c-erbB-2 two single chain Fv construct.
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CREATIVE BIOMOLECULES INC.
                              ISSSSGSSSSSSSSSSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT 176
                                                                                 ADDFKERFAPSLETSATTAHLQINNLRNEDSATYFCARRF----GFAYWGQGTLVSVSAS
                                                                                                                                                   EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                                                AEEFKGRPAFSLETSASTAYLQINNLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA-
                                                                                                                             EIQLVQSGPELKKPGETVKISCKASGYTFTNYGMNWVKQAPGKGLKWM;3WINTNTGEPTY
                                                                                                                                                                                                                                                            250 AA;
- SSSSGSSSSGSSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAWYQQKPGQS
                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ring DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9208-00831967.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9308-00133804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "claimed C-terminal tail to facilitate crosslinking of two sFv polypeptides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    246. .250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label=
                                                                                                                                                                                                            72.7%; Score 914; DB 2; 71.7%; Pred. No. 1.4e-61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Huston JS,
                                                                                                                                                                                             32;
                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Houston
                                                                                                                                                                                                                          DB 2; Length 250;
                                                                                                                                                                                             30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genes of the 741F8 anti-c
d from the cDNA of the
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RESULT 11
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ID AW29
XX AW29
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XX Adenc
KW Adenc
KW Expre
KW Itrans
KW Bingl
XX Syntl
XX US56E
PN US56E
XX US56E
XX US56E
XX UF1
DR N-FSI
XX PPT
DR N-FSI
XX PPT
DR N-FSI
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                                                                                        This sequence is a single chain anti-c-erbB-2 antibody (scFv) derived CC from hybridoma 741F8. The protein product includes a C-terminal tail Gly4 CC -Cys in some constructs. Single-chain antibodies can be produced using a CC novel method which comprises culturing an immmortalised eukaryotic cell containing transfected DNA sequences (encoding the protein of interest). CC integrated into its genome. In particular expression effector vectors containing a non-native reporter DNA (encoding the scFv) and viral CC sequences to promote transcription and translation (e.g. the adenovirus CC ElA and VA1 genes as shown in AAT91831 and AAT91834 respectively) are used. The scFv that is produced, when properly folded, has a structure CC with mono- or bi-functional binding activity. The method is especially CC used to produce single-chain antibody fragments (scFv), e.g. for imaging CC tumours or delivering therapeutic agents to them, particularly breast and CC contain cancers that express the c-erbB-2 antigen. Other scFv are used in model studies and for treating digoxin intoxication. (Updated on 25-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Producing single chain binding protein in immortalised eukaryotic cells which comprise protein coding sequences, a transcription activator and translation promotion sequences, provides high expression at low copy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Adenovirus; ElA; transactivator; transcription activator; stimulate; expression vector; single-chain binding protein; VAI; enhance; PCR; translation; production; immortal; eukaryotic cell; scFv; primer; single-chain antibody fragment; imaging; tumour; breast cancer; ovarian cancer; c-erbB-2 antigen; digoxin intoxication.
       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Col 29-32; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oppermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
07-JAN-1998
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                                                                      to correct
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Query Match Best Local S

Best Local Similarity Matches 177; Conserv

Conservative

72.7%; Score 914; 71.7%; Pred. No. 1 32;

Mismatches

Indels

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Gaps

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1.4e-61;

DB 2; Length 250;

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chain binding protein, especially a single chain antibody, by yellocation binding protein, especially a single chain antibody, by yellocation activator protein that acts on and stimulates a viral promoter controlling the expression of DNA encoding the single-chain binding protein, such as the adenovirus Ad2 ELA protein encoded by the sequence AAT78873, or an RNA sequence able to promote translation of the RNA transcript from the heterologous gene, such as the adenoviral VA1 gene (AAT78876). The sequence presented here is the amino acid sequence of the anti-c-erbB2 single chain antibody 741. The coding sequence was constructed by amplifying the variable heavy and light chain genes which were then cloned into a pUC vector. For secretion from mammalian cells, the sequence is preceded by the signal peptide sequence from the
                                                                                                                                                                      The invention relates to methods of increasing production of a single-chain binding protein, especially a single chain antibody, by generati
                                                                                                                                                                                                                                                         Production of cell line for producing single-chain binding protein using construct containing DNA encoding viral transcription activates
                                                                                                                                                                                                                                                                                                                                                   Oppermann
                                                                                                                                                                                                                                                                                                                                                                                                             25-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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09-OCT-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single chain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ISSSSGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABEFKGRPAFSLETSASTAYLQINNLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
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                                                                                                                                                                                                                    Col 29-32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chain reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single-chain, binding protein, antibody, eukaryote,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibody 741F8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
                                                                                                                                                                                                                                                                                                                                                   Dorai
                                                                                                                                                                                                                                                                                                                                                                                                           93US-00143498
                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-00461184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ivator; promoter; expression; adenovirus; EIA; PCR;
reaction; amplification; primer; herpes simplex virus;
vector; enhancer; translation; heterologous.
                                                                                                                                                                                                                 24pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAMYQQKPGQS
                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              250
                                                                                                                                                                                                                                                                                                                                                                                INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B
                                                                                                                                                                                                                                                             activator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           238
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RESULT 13
AAW47012
ID AAW47
XX AAW47
XX AAW47
AC AAW47
AC AAW47
XX Singl
XX Singl
XX Singl
XX US573
XX
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អនុស្ន
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Sim
Matches 177;
                                                                                                                                                                                                                                           Immortalised eukaryotic cell comprising transfected DNa sequences useful for enhanced production of proteins encoded by non native (
                                                                                                                                                                                                                                                                                                                                                                                                       Oppermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US5733782-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein production; non-native gene; hard to express gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transactivating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Single chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW47012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW47012 standard; protein; 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CREA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 250 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         monoclonal antibody 520C9 (AAT78880) 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                     1998-229831/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         CREATIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LEIKRAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRF----GFAYWGQGTLVSVSAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EIQLVQSGDELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LEIKRAD 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IKRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ISSSSGSSSSSSSSSSSIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEEFKGRFAFSLETSASTAYLOINNLKNEDTATYFCGROFITYGGFANWGQGTLVTVSA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EIQLVQSGPELKKPGETVKISCKASGYTFTNYGMNWVKQAPGKGLKWMGWINTNTGEPTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PKLLIYWTSTRHTGVPDRFTGSGSGTDYTLTISSVQAEDLALHYCQQHYRVPYTFGGGTK 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -SSSSSSSSSSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAWYQQKPGQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 binding site molecule 7414F8; sFv 741F8; large sing transcription activator; viral transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              binding site molecule
                                                                                                                                                                                                                                                                                                                                                                                                       Dorai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93US-00143498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-00464589
                                                                                                                                                                                                                                                                                                                                                                                                                                                         BIOMOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            72.7%;
71.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 914; DB 2;
Pred. No. 1.4e-61;
2; Mismatches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (sFv) 741F8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or
Or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                             genes
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The present sequence represents single chain binding site molecule (sFv). The sequence was expressed to exemplify the invention, which is a method for the expression of poorly expressed genes in an immortalized eukaryotic cell. This cell contains transfected DNA sequences operatively integrated into its genome. The transfected DNA sequences encode a viral transcription promoter linked to a non-native reporter sequence encoding a single chain binding protein. The promoter is activated by a viral

operatively

Example

2

Col 31-32; 24pp;

English.

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RESULT 14
AAW53168
ID AAW53
XX AAW53
XX AAW53
AC AAW53
XX AAT19
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 177; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW53168 standard; protein; 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transcription activator protein. The cell allows the production on a commercial scale of proteins encoded by non-native hard to express genes
     Example 1; Col 25-28;
                                                                          Imaging of antigens in
                                                                                                                             N-PSDB; AAV21796.
                                                                                                                                                 WPI; 1998-311318/27.
                                                                                                                                                                                                  Oppermann H,
                                                                                                                                                                                                                                                                                                                          06-FEB-1992;
07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                  05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour; diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antigen imaging; single chain Fv; sFv; linker; dimeric; cancer; c-erbB-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              741F8 anti-c-erbB-2 sFv' dimeric construct protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JUL-1998 (first entry)
                                                                                                                                                                                                                                              (CHIR ) CHIRON CORP.
(CREA-) CREATIVE BIOMOLECULES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LEIKRAD 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PKLLIYWTSTRHTGVPDRFTGSGSGTDYTLTISSVQAEDLALHYCQQHYXVPYTFGGGTK 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IKRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYA[FPYTFGGGTN 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -SSSGSSSGGSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAMYQQKPGQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRF----GFAYW3QGTLVSVSAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LEIKRAD 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ISSSSSSSSSSSSSSSSIOMTOSPSSLSASIGERVSLTCRASODIGNSLTWLOOEPDGT
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                                                                                                                                                                                             Ring DB,
                                                                                                                                                                                                                                                                                                                        92US-00831967.
93US-00133804.
                                                                                                                                                                                                                                                                                                                                                                                                  95US-00461838
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        246. .250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "linker peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "Gly4-Cys C-terminal tail"
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30pp; English
                                                                            VÍVO
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Pred. No. 1.4e-61;
2; Mismatches 30
                                                                            dimers
                                                                                                                                                                                                  Houston
                                                                     of single-chain antibody
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This represents the protein sequence of a 741F8 sFv' (single chain Fv) C terminal Gly4-Cys construct. This was constructed by connecting the Vh and Vl genes with a DNA sequence encoding a 14 residue polypeptide linker. 741F8 is a monoclonal antibody useful in targeting c-erbB-2 antigen. This dimeric construct can be used in the methods of invention of imaging a preselected antigen expressed in a mammal. The methods are used in magnetic resonance imaging of c-erbB-2 or related antigens in cancer diagnosis. The biosynthetic constructs have enhanced properties a in vivo targetting agents in comparison with intact monoclonal antibodie or their Fab fragments. The dimeric constructs permit the in vivo targetting of an epitope on an antigen with greater apparent avidity, including greater tumour specificity, tumour localisation and tumour retention properties than that of the Fab fragment having the same CDRS
                                                                                                                                                                                      Key
Peptide
                                                                                                                                                                                                                                                                                                                                Anti-c-erbB-2 sFv' protein sequence
                                                                                                                                                                                                                                                                                                                                                                                             AAW80422;
                                06-FEB-1992;
07-OCT-1993;
                                                                              05-JUN-1995;
                                                                                                                                            US5837846-A.
                                                                                                                                                                                                                                                                                   targeted delivery; antigen-expressing cell.
                                                                                                                                                                                                                                                                                                Anti-c-erbB-2 sFv'; c-erbB-2; antigen; tumour cell; antibody 741F8;
                                                                                                                                                                                                                                                                                                                                                               28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                          AAW80422 standard; protein;
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                                                                                                             17-NOV-1998
                                                                                                                                                                                                                                     Homo sapiens.
   (CREA-) CREATIVE BIOMOLECULES INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ISSSGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEEFKGRFAFSLETSASTAYLQINNLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA- 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRF----GFAYWGQGTLVSVSAS 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LEIKRAD 245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -ŚŚŚŚŚŚŚŚŚŚŚŚŚŚ--ŚDIVMTQŚPKFMŚTŚVĆDRVŚISCKAŚQDVSTAVAWYQQKPGQS
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                                92US-00831967.
93US-00133804.
                                                                              95US-00461386
                                                                                                                                                                                        Location/Qualifiers 122. .135
                                                                                                                                                                           'note= "linker"
                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72.7%; Score 914; DB 2;
71.7%; Pred. No. 1.4e-61;
tive 32; Mismatches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Length 250;
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Search completed: April 4,
Job time : 488.224 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents an anti-c-erbB-2 sFv'. c-erbB-2 is an antigen that is overexpressed on the surface of tumour cells. An antibody designated 741F78 binds c-erbB-2. Variable heavy and light sequences of antibody 741F8 are connected, together with a linker, to produce the present single chain Fv gene. Anti-c-erbB-2 sFv' exemplifies the invention. Dimers of the single chain Fv are used for targeted delivery of drugs or imaging agents (e.g. cytotoxins, prodrugs or 99m-technetium) to antigen-expressing cells, particularly for treatment or diagnosis of tumours (especially of ovary or breast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid encoding single-chain Fv fragment specific for antigens - and having C-terminal tail for crosslinking to form dimer with improved pharmacokinetic properties, used to deliver drugs and imaging agents, especially to tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 250 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                                                                                                                                                                                                                                                                                                                            -SSSSGSSSSGSSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAWYQQKPGQS
                                                                                                               LEIKRAD 245
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Result
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Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
          110
110
111
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113
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                                                                                                                                                           1258
1063.5
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902
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Gapop 10.0 ,
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77.6
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1258
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Copyright (c) 1993 - 2005 Compugen Ltd.
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/cgn2_6/ptodata/1/pubpaa/US09B_UBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US10REW_PUB.pep:*
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/ Cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
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/cgn2_6/ptodata/1/pubpaa/PCT
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US-09-766-543-10
US-09-766-543-10
US-09-766-543-10
US-09-766-543-2
US-10-897-853-2
US-10-893-94-6
US-10-879-94-6
US-10-879-94-6
US-09-924-099-0
US-09-924-099-10
US-09-10-177-243-148
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Sequence 6, Appli
Sequence 6, Appli
Sequence 12, Appl
Sequence 13, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 10, Appl
Sequence 148, App
Sequence 148, App
Sequence 148, App
Sequence 148, App
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e 28,	Sequ	0-29	16	899	8	735	
e 7,	æ	-10-496-179-	17	896	8	735	
e 3,	Ð	-10-496-17	17	896	8	735	ü
e 2,	æ	-10-496-179-	17	896	8	735	
e 26,	Segn	0-296-085A-2	16	896		735	14
e 18,	Segn	-10-296-08	16	896	8	735	
e 17,	Sequ	-10-296-085A-1	16	896	8	735	
4,	•	-10-496-17	17	895	8	735	
		US-10-296-085A-27	16	895	в.	735	
20,		-10-296-085A-2	16	895	8	735	
e 19,	Seqn	-10-296-085A-1	16	895	8.	735	35
e 932	Sequ	0-293-418-93	15	250	8	740.5	
e 932	Seqn	-09-880-74	10	250	8	740.5	
e 17:	æ	-10-293-418-177	15	248	9	743.5	32
e 177	O	-09-880-748-177	10	248	9	743.5	
e 144	Seqn	-10-293-418-1	15	248	0	762.5	
e 14	Seqn	9-880-748-	10	248		762.5	
e 75,	Segn	-10-239-656-	15	503		764	
e 193	Segn	-10-293-418-1	15	251	0	764	
e 193	Seqn	-09-880-748-	10	251	٥.	764	
e 11(Segn	-10-293-418-	15	248		9	
e 110	Segn	-09-880-748-	10	248		766.5	
e 10	Segn	0-071-485-1	14	230		9	
e 93,	Segn	-10-071-485-9	14	235		779	
e 77,	Segn	-10-239-656	15	503	'n	~1	
e 37,	Sequ	-10-222-026A-	14	287	Ņ	782.5	
e 79,	Sequ	0-239-656-7	15	505	٧.	786	
e 91,	Sequ	-10-071-485	14	240	'n	791.5	
e 90,	Sequ	0-071-485-9	14	711	ω	ö	
e 85,	Segu	0-071-485-8	14	541	'n	Ö	
e 2	0	-10-071-	14	267	63.9	804	5
quence 67, Appl	Ø	0-239-656	15	ū			

ALIGNMENTS

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US-09-887-853-6
US-09-887-853-6
J Sequence 6, Application US/09887853
Patent No. US20020168375A1
Patent No. US20020168375A1
PATENDAL INFORMATION:
Ring, David B.
RITTLE OF INVENTION: Biosynthetic Binding Proteins For INVENTION: ADDRESSE: Testa, Hurwitz & Thibeault/Patent Department STREET: Exchange place, 53 State Street CITY: Boston STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/09/887,853
FILING DATE: 21-Unn-2001
CLASSIFICATION DATA:
APPLICATION DATA:
APPLI
```

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GENERAL INFORMATION:

APPLICANT: HUSEON, J.

APPLICANT: HOUSEON, L.L.

APPLICANT: Ring, D.

APPLICANT: Ring, D.

APPLICANT: Oppermann, H.

ITILE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING

FILE REFERENCE: CIBT-P01-130

CURRENT APPLICATION NUMBER: US/09/683,547

CURRENT FILING DATE: 2003-10-10

PRIOR APPLICATION NUMBER: US/09/558,741

PRIOR APPLICATION NUMBER: 07/831,967

PRIOR APPLICATION NUMBER: 07/831,967

PRIOR FILING DATE: 1992-02-06

INMBER: OF SEQ ID NOS: 16

SOFTWARE: PatentIn version 3.1

SEQ ID NO 6

LENGTH: 243
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US-10-683-547-6
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                                                                                                                                Query Match
Best Local Similarity
Matches 243; Conser
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                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                  FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 100.0%; Score 1258; DB 9; Local Similarity 100.0%; Pred. No. 7.8e-86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   241 RAD 243
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ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS 120
                                                         EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                                                                EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY 60
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                                                                                                                                100.0%; Score 1258; DB 17; ilarity 100.0%; Pred. No. 7.8e-86; Conservative 0; Mismatches 0;
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RESULT 4
US-10-239-656-73
US-10-239-656-73
Fequence 73, Application US/10239656
Publication No. US20040038339A1
TYPOPMATION:
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Patent No. US20020041865A1
GENERAL INFORMATION:
APPLICANT: Austin, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Kwok, Cheuk S.
APPLICANT: Kwok, Cheuk S.
FITLE OF INVENTION: METHODS FOR TREATING TUMORS
FILE REFERENCE: PP01679.002
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 12
LENGTH: 276
TYPE: PRT
ORGANISM: Artificial Sequence
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OTHER INFORMATION: Description
OTHER INFORMATION: linker
US-09-766-543-12
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US-09-766-543-12
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Local Similarity 83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 1063.5; DB Pred. No. 2.6e-71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of Artificial Sequence: H520C9sFv
                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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APPLICANT: KUPER, PETER
APPLICANT: KIETHMULLER, GERT
APPLICANT: LUTTERBUSE, RALF
APPLICANT: BORSCHERT, KATRIN
APPLICANT: KISCHEL, ROMAN
APPLICANT: MAYER, MONIKA
APPLICANT: HOFMEISTER, ROBERT

```
GENERAL INFORMATION:
APPLICANT: AUSLIN, Richard
APPLICANT: Kwok, Cheuk S.
APPLICANT: Kwok, Cheuk S.
APPLICANT: King, David B.
TITLE OF INVENTION: METHODS FOR TREATING TUMORS
PILE REFERENCE: PP01679.002
CURRENT APPLICATION NUMBER: US/09/766,543
CURRENT FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR APPLICATION NUMBER: 60/177,258
PRIOR APPLICATION NUMBER: 2000-01-20
NUMBER OF SEG ID NOS: 14
SOPTWARE: Patentin Ver. 2.0
SEG ID NO 10
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CURRENT FILING DATE: 2003-03-06
PRIOR APPLICATION NUMBER: PCT/EP01/03414
PRIOR FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: EP 00106467.4
PRIOR PILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 92
SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 73
SEQ ID NO 73
                                                                                                                                                                                                                                                                                                                 RESULT 5
US-09-766-543-10
18-09-766-543-10
18-09-766-543-10
18-09-766-543
18-09-766-543
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: MULTIFUNCTIONAL POLYPEPTIDES COMPRISING A BINDING SITE TITLE OF INVENTION: TO AN EPITOPE OF THE NKG2D RECEPTOR COMPLEX FILE REFERENCE: 029976/0106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Synthetic 3B10xP4-OTHER INFORMATION: 3 bispecific single chain Fv
ORGANISM: Artificial Sequence
                                      ENGTH: 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 77.6%; Score 976; DB 15; Local Similarity 55.1%; Pred. No. 1.6e-64;
                                                                                                                                                                                                                                                                                                                                                                                                                                      488
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        248 EVQLLESGGGLVQPGGSLKLSCAASGFDFSRYWMSWVRQAPGKGLEWIGEINPDSSTINY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
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                                                                                                                                                                                                                                                                                                                                                                                                                                    R 488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYIFGGGTNLEIK 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRL 180
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RESULT 6
US-09-887-853-2
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Patent No. US20020168375A1
GENERAL INFORMATION:
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Best Local
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                                               INFORMATION FOR SEQ ID NO: 2:
                                                        PRIOR PETENCHION NUMBER: US/09/887,853
PILING DATE: 21-Jun-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US/08/133,804
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7100
                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PAtentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                       ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF
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Similarity 71.6%; Pred. No. 6.4e-63;
74; Conservative 32; Mismatches 34
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STREET: Exchange Place,
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                                                                                                                                                                                                                                                                                                                                                                                                            STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Boston
LENGTH: 250 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ring, David B.
INVENTION: Biosynthetic Binding Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Huston, James S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oppermann, Hermann
Houston, L. L.
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                                                                                                                                                                                                                                                                                              Version
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Sequence 2, Application US/10683547

Publication No. US20050058638A1

GENERAL INFORMATION:
APPLICANT: Huston, J.
APPLICANT: Houston, L.L.
APPLICANT: General, H.
APPLICANT: Oppermann, H.
APPLICANT: Oppermann, H.
APPLICANT: Oppermann, H.
CITILE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING
FILE REFERENCE; CIBT-P01-130

CURRENT APPLICATION NUMBER: US/10/683,547

CURRENT FILING DATE: 2003-10-10
PRIOR APPLICATION NUMBER: US/09/558,741

PRIOR APPLICATION NUMBER: US/09/558,741

PRIOR APPLICATION NUMBER: 07/831,967

PRIOR FILING DATE: 1992-02-06

NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1

SEQ ID NO
LENGTH: 250
TYPE: PRT
ORGANISM: Artificial Sequence
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; MOLECULE TYPE: protein

; SEQUENCE DESCRIPTION: SEQ ID

US-09-887-853-2
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                                                                                                                                                                                                                                                                     Query Match
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: 741F8 sFv'
                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          117 ISSSSGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT 176
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                                                                                                                                                                               1 EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWKKQAPGKGLKWMGWINTYTGQSTY
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                                                                                                             ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRF----GFAYWGQGTLVSVSAS
                                     ISSSSGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT 176
                                                                           AEEFKGRFAFSLETSASTAYLQIKKLKNEDTATYFCGRQFITYGGFANWGQGTLVTVSA-
                                                                                                                                                     BIQLVQSGPELKKPGETVKISCKASGYTFTNYGMNWVKQAPGKGLKWMGWINTNTGEPTY
-SSSGSSSGSSS--SDIVMTQSPKFMSTSVGDRVSISCKASQDVSTAVAWYQQKPGQS
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71.7%;
                                                                                                                                                                                                                                                 71.7%; Score 902; DB 17; 70.9%; Pred. No. 2.4e-59;
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; Pred. No. 3.1e-60;
32; Mismatches 30
                                                                                                                                                                                                                               32; Mismatches
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                                                                                                                                                                                                                                                                   Length 250;
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APPLICANT: Karow, Margaret

APPLICANT: Smith, Eric

APPLICANT: Smith, Eric

APPLICANT: Smith, Eric

APPLICANT: Smith, Eric

FITLE OF INVENTION: USE

FILE REPRENCE: REG 203E2

CURRENT APPLICATION NUMBER: US/10/879,994

CURRENT FILING DATE: 2004-06-29

PRIOR APPLICATION NUMBER: 10/610,452

PRIOR APPLICATION NUMBER: 10/610,452

PRIOR FILING DATE: 2003-06-30

NUMBER OF SEQ ID NOS: 112

SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                          Sequence 9, Application US/09924099 Patent No. US20020128450A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 168; Conservative
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Publication No. US20050032175A1
APPLICANT: NISHIDA, Yoshihiro
APPLICANT: OKURA, Takanori
APPLICANT: TANIMOTO, Tadao
APPLICANT: TANIMOTO, Tadao
APPLICANT: KURIMOTO, Masashi
TITLE OF INVENTION: PEPTIDE
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/09/924,099
CURRENT FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/338,511
PRIOR FILING DATE: EARLIER FILING DATE: 1999-06-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 236
                                                                                                                                                                                                                                                                                                                                                                                      236 R
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 NQKFRDKATLTVDQSSTTAFMHLNSLTSEDSAVYFCAR--GLRFWGQGTLVTVSA---GG 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARREGEAYWGQGTLVSVSASISSS 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 68.5%; Score 861.5; DB 1 Similarity 69.7%; Pred. No. 2.3e-56; 68; Conservative 26; Mismatches 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNLEIK 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SGSSSSGSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVDYYCLQYASSPYTFGGGTKLAIK 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGSGGGGGGGGDIQMTQSPSSLSASLGERVSLTCRASQDIGSKLYWLQQEPDGTFKRL 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EIQLQQSGPELVKPGASVKVSCKASGYSFTDYFIYMVKQSHGKSLEWIGDIDPYNGDTSY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LEIKRAD 245
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US-09-924-099-10
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CURRENT FILING DATE: 2001-08-08

PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/338,511

PRIOR FILING DATE: EARLIER FILING DATE: 1999-06-23

PRIOR FILING DATE: EARLIER FILING DATE: 1999-06-23

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: JP 289,044,

PRIOR FILING DATE: EARLIER FILING DATE: 1998-10-12

PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-22

NUMBER OF SEQ ID NOS: 33

SEQ ID NO 10

LENGTH: 243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10, Application US/09924099 Patent No. US20020128450A1
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Best Local
                                                                                      Matches
                                                                                                                              Query Match
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                                                                                                                                                                            -09-924-099-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: NISHIDA, YOShihiro
APPLICANT: OKURA, Takanori
APPLICANT: TANIMOTO, Tadao
APPLICANT: KURIMOTO, Masashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: TANIMOTO, Tadao
APPLICANT: KURIMOTO, Masashi
TITLE OF INVENTION: PEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE:
                                                                                                                                                                                              OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Artificially produced peptide in the form of a single OTHER INFORMATION: chain
OTHER INFORMATION: variable region fragment (scFv) which neutralizes IL-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 237
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                FEATURE:
                                                                                   Local Similarity
nes 167; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 69.6
nes 167; Conservative
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                                    EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRL 180
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EIQLQQSGPELVKPGASVKVSCKASGYSFTDYFIYWVKQSHGKSLEWIGDIDPYNGDTSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVDYYCLQYASSPYTFGGGTKLAIK
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                                                                                      Conservative
                                                                                                                                                                                         Artificially produced peptide in the form of a single chain variable region fragment (scFv) which neutralizes IL-18
                                                                                                        69.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68.2%; Score 857.5; DB 9; 69.6%; Pred. No. 4.7e-56;
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                                                                                   26;
                                                                                 Score 857.5; DB 9;
Pred. No. 4.8e-56;
26; Mismatches 44;
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                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: JP 365,023/98
DATE: 1998-12-22
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                                                                                   Indels
                                                                                                                            Length
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US-10-127-890-148
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                                                        LENGTH: 240 amino acids
TYPE: amino acid
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
3-10-127-890-148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 148, Application US/10 Publication No. US20030166196A1
Query Match
Best Local Similarity
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                                                                                                                                                                                                                   TELEX: 650 388-1248 INFORMATION FOR SEQ ID NO: 14
                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/646,360
FILING DATE: 13-MAY-1996
APPLICATION NUMBER: PCT/US94/05348
FILING DATE: 12-MAY-1994
APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-1993
APPLICATION NUMBER: US 07/988,430
FILING DATE: 09-DEC-1992
APPLICATION NUMBER: US 07/901,707
FILING DATE: 19-UN-1992
APPLICATION NUMBER: US 07/787,567
FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Better, Marc D.
Carroll, Stephen F.
Studnika, Gary M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: McAndrews, Held & Malloy, Ltd.
STREET: 500 West Madison Street, 34th floor
                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-8889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Studnika, Gary M.
TITLE OF INVENTION: Immunotoxins Comprising Ribosome-Inactivating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         181 IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYYCLQYAIFPYTFGGGTNLEIK 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFAYWGQGTLVSVSASISSS 120
                                                                                                                                                                                                                                                                                                                                                 NAME: McNicholas, Janet M. REGISTRATION NUMBER: 32,918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/10/127,890 FILING DATE: 23-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVDYYCLQYASSPYTFGGGTKLAIK
                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 200-70.P4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 60661
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                                                                                                                                                                                                                                                              TELEFAX:
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                                                                                                                                                                                                                                                                312/707-9155
                                                                                                                                                                                                                     ID NO: 148:
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  65.7%;
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    Score 827; DB 14;
Pred. No. 8.8e-54;
                                                                                     148:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Version
                      Length 240;
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RESULT 12
US-10-717-243-148
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Publication No. US2005
GENERAL INFORMATION:
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                  INFORMATION
                                                                                ATTORNEY/AGENT INFORMATION:
NAME: MCNicholas, Janet M.
REGISTRATION NUMBER: 32,918
REFERENCE/DOCKET NUMBER: 11022US09/200-70.P3.C3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-8889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: McAndrews, Held & Malloy, Ltd.
STREET: 500 West Madison Street, 34th floor
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Immunotoxins Comprising Ribosome-Inactivating Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Better, Marc D. Carroll, Stephen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        178
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                                                                                                                                                                                                             APPLICATION NUMBER: US/08/839,765
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 08/425,336
FILING DATE: 18-APR-1995
APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-193
APPLICATION NUMBER: US 07/988,430
FILING DATE: 09-DEC-1992
APPLICATION NUMBER: US 07/901,707
FILING DATE: 09-DEC-1992
APPLICATION NUMBER: US 07/901,707
FILING DATE: 09-UN-1992
APPLICATION NUMBER: US 07/787,567
FILING DATE: 04-NOV-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 18-Nov-2003
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Chicago
STATE: Illinois
COUNTRY: USA
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                  TELEX: 650 388-1;
N FOR SEQ ID NO:
                                                              TELEFAX: 312/707-9155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/10/717,243
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CHARACTERISTICS
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MOLECULE TYPE: protein
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SEQUENCE DESCRIPTION: SEQ ID
US-10-717-243-148
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                                                                                                              Matches
                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                              SEQ ID NO 55
LENGTH: 252
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: KISCHEL, ROMAN
APPLICANT: MAYER, MONIKA
APPLICANT: HOFMEISTER, ROBERT
TITLE OF INVENTION: MULTIFUNCTIONAL POLYPEPTIDES COMPRISING A BINDING SITE
TITLE OF INVENTION: TO AN EPITOPE OF THE NKG2D RECEPTOR COMPLEX
FILE REFERENCE: 0.2976/0106
CURRENT FILINGTION NUMBER: US/10/239,656
CURRENT FILING DATE: 2003-03-06
PRIOR APPLICATION NUMBER: PCT/EP01/03414
PRIOR APPLICATION NUMBER: PCT/EP01/03414
PRIOR FILING DATE: 2001-03-26
                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 92
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: EP 00106467.4
                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: chain Fv
                                                                                                              160;
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61 ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARR---FGFAYWGQGTLVSVSASI 117
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LUTTERBUSE, RALF
BORSCHERT, KATRIN
KISCHEL, ROMAN
                                                                     EIQLVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTY
                                  EVQLLESGGGLVQPGGSLKLSCAASGFDFSRYWMSWVRQAPGKGLEWIGEINPDSSTINY
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                                                                                                              Conservative
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                                                                                                                           65.0%; Score 818; DB 15; 65.6%; Pred. No. 4.3e-53;
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Pred. No. 8.8e-54;
Pred. No. 8.8e-54;
                                                                                                              Mismatches
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                                                                                                                                           Length
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                                                                                                          Gaps
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APPLICANT: KISCHEL, ROMAN

APPLICANT: MAYER, MONIKA

APPLICANT: MAYER, MONIKA

APPLICANT: MOPMEISTER, ROBERT

TITLE OF INVENTION: MULTIFUNCTIONAL POLYPEPTIDES COMPRISING A BINDING SITE

TITLE OF INVENTION: TO AN EPITOPE OF THE NKG2D RECEPTOR COMPLEX

FILE REFERENCE: 029976/0106

CURRENT APPLICATION NUMBER: US/10/239,656

CURRENT FILING DATE: 2003-03-06

PRIOR APPLICATION NUMBER: POT/EP01/03414

PRIOR APPLICATION NUMBER: EP 00106467.4

PRIOR APPLICATION NUMBER: EP 00106467.4

PRIOR FILING DATE: 2000-03-24

NUMBER OF SEQ ID NOS: 92

SOFTWARE: PATENTIN Ver. 2.1

SEQ ID NO 67

LENGTH: 257
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic P5-10 single ; OTHER INFORMATION: chain Fv
US-10-239-656-67
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US-10-239-656-67
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APPLICANT: KIETHMULLER, GERT
APPLICANT: KIETHMULLER, RALF
APPLICANT: BORSCHERT, KATRIN
APPLICANT: KISCHEL, ROMAN
APPLICANT: MAYER, MONIKA
APPLICANT: HOFMEISTER, ROBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 67, Application US/10239656 Publication No. US20040038339A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               Matches 161;
                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                              Match 64.7%; Score 813.5; DB 15; Length 257; Local Similarity 64.7%; Pred. No. 9.5e-53;
 238
                                   233
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                                                                                                                                                                                     VSASISSSGSSSSGSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQE 172
                                                                                                                                                                                                                      YDPKFQGKASITADTSSNTAYLQLSSLTSBDTAAYYCASFYYYGSSYRYFDVMGQGTTVT
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GGTKLEIKR 246
                                                                                                 PDGTIKRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVVYYCI,QYAIFPYTFG
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                                   GGTNLEIKR 241
                                                                          PDGTIKRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVDYYCI,QYASSPYTFG
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Search completed: April Job time: 359.677 secs

4, 2005, 16:42:42

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GENERAL INFORMATION.

APPLICANT: Buyes, Marie-Ange
APPLICANT: Sablon, Erwin
TITLE OF INVENTION: INTERFERON-gamma-BINDING MOLECULES FOR TREATING SEPTIC
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: SHOCK,
TITLE OF INVENTION: CACHEXIA, IMMUNE DISEASES AND SKIN DISORDERS
FILE REFERENCE: INNS.015
CURRENT APPLICATION NUMBER: US/10/071,485
CURRENT FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 09/485,737
PRIOR APPLICATION NUMBER: DEC/EP 98/05165
PRIOR APPLICATION NUMBER: PCT/EP 98/05165
PRIOR APPLICATION NUMBER: EPO 98870139.7
PRIOR FILING DATE: 1998-06-18
PRIOR FILING DATE: 1997-08-18
NUMBER OF SEQ ID NOS: 104
SOFTWARE: Patentin version 3.0
SEQ ID NO 2
1.FNCTH: 267
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Best Local S
Matches 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 267
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                            179
                                                                                                                                                                                     140 GGGGGGGGGGGDIVLTQSPATMSASPGERVTLTCSASSSI-SYMFWYHQRPGQSPR 198
259
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                                                                                          199 LLIYDTSNLASGVPARFSGSGSGTSYSLTISRMEPEDFATYFCHQSSSYPFTFGQGTKLE
                                                                                                                                                                                                              119 SSSGSSSSGSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIK 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tch 63.9%; Score 804; DB 1 al Similarity 63.8%; Pred. No. 5e-52; 155; Conservative 32; Mismatches
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Result
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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single chain Fv an
Ig heavy chain pre
L6 mab heavy chain
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single chain Fv antibody - mouse (;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_cha C;Accession: S41374
C;Accession: S41

06-Jan-1995 #text_change 06-Jan-1995

of a

single chain

Fv antibody

A; Reference number: S41374

RESULT S41374

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179

KLWVYYTSHLPPGVPARFSGSGSGNSYSLTISSMEGEDAATYYCQQFTSSPFTFGSGTKL KRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNL SGGGGSGGGDSGGGGSDIELTQSPAIMSASLGEKVTMSCRASSSV-NFIYMYQQKSDASP

178 120 118 63

62

SSSGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTI

178 177

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Query Match Best Local Matches 11	RESULT 1 A56446 C; Species C; Species C; Access R; Tang J; Biol. A; Title: A; Title: A; Referse A; Access		8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
Query Match Best Local Similarity Matches 117; Conser	RESULT 1 A56446 Ig heavy chain V region C; Species: Mus musculus C; Date: 19-Jan-1996 #sec C; Accession: A56446 R; Tang, P.M.; Foltz, L.; J. Biol. Chem. 270, 782; A; Title: A high affinit; A; Reterence number: A5646 A; Accession: A56446 A; Accession: A56446 A; Actus: preliminary A; Molecule type: mRNA A; Residues: 1-268 < TANA A; Residues: 1-268 < TANA A; Residues: heterotetran		432 4334 42333 4323 418.5 418 418 415 415 405 406
47. ilarity 47. Conservative	RESULT 1 A56446 Ig heavy chain V region (3H-3H scFv) - mou C;Species: Mus musculus (house mouse) C;Species: Mus musculus (house mouse) C;Late: 19-Jan-1996 #sequence_revision 19-C;Accession: A56446 C;Accession: A56446 J. Baiol. Chem. 270, 7829-7835, 1995 A;Title: A high affinity digoxin-binding ph;Reference number: A56446; MUID:95229583; A;Reference number: A56446; MUID:95229583; A;Accession: A56446 A;Battus: preliminary A;Cross-references: GB:UZ0617 C;Keywords: heterotetramer; immunoglobulin		60000000000000000000000000000000000000
47.0%; 47.8%; ative	gion (3H-3H scFv) ulus (house mouse the guerne_revisi L.A.; Mahoney, 78.9-7835, 1995 Sinity digoxin-bin A56446; MUID:952 TAN GB:U20617 GB:U20617		108 105 105 105 105 105 105 114 105 114 115 115 115
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2; 75;	(strain BALB/C) n-1996 #text_cheneler, P.A. nueler, P.A. tein displayed of MID:7713873		
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L6 mAb heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Bate: 27-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: B42848; S33903
R;Fell, H.P.; Gayle, M.A.; Yelton, D.; Lipsich, L.; Schieven, G.L.; Marken,
J. Biol. Chem. 267, 15552-1558, 1992
A;Title: Chimeric L6 anti-tumor antibody. Genomic construction, expression,
A;Reference number: A42848; MUID:92348410; PMID:1639794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: synthetic C;Date: 13-Jan-1995 #sequence_revision 30-Apr-1998 #text_change 20-Oct-2000 C;Date: 13-Jan-1995 #sequence_revision 30-Apr-1998 #text_change 20-Oct-2000 C;Accession: S33905 R;Liu, A.Y.; Robinson, R.R.; Hellstroem, K.E.; Murray Jr., E.D.; Chang, C.P.; Hells Proc. Natl. Acad. Sci. U.S.A. 84, 3439-3443, 1987 A;Title: Chimeric mouse-human IgG1 antibody that can mediate lysis of cancer cells. A;Reference number: S33905; MUID:87204152; PMID:3106970 A;Accession: S33905
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A; Residues: 1-146 < LIU>
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-249 <ART>
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                                                                                                                                                                                                                                                                                                                                            ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFGFA-----YWGQGTLVSVS
                                                                                                                                                                                                                                                                                                                                                                                                 QIQLVQSGPELKKPGETVKISCKASGYTFTNYGMNWVKQAPGKGLKWMGWINTYTGQPTY 79
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                                                                                                                                                                                                                                                                              AS 116
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                                                                                                                                                                                                                                           SA 140
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Pred. No. 1.5e-30;
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Mismatches
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                            and
                                                                J.S.; Aruffo
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R;Froyen, G.F.V.
submitted to the EMBL Data Library, May 1993
A;Reference number: S35759
A;Accession: S35759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EAUVULU protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Aate: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 21-Jan-2000
C;Aacesslon: S35759
R;Froyen, G.F.V.
A;Accession: S32190
A;Status: preliminary
A;Molecule type: mRNA
                                                                                                  Ig heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 21-Jan-2000
C;Accession: S32190
                                                                                                                                                                        RESULT
S32190
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C;Superfamily: immunoglobulin V region;
F;15-98/Domain: immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues: 1-136 <FRO>
A;Cross-references: EMBL:X72796; NID:g312496; PIDN:CAA51316.1; PID:g312497
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;34-117/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Accession: B42848
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-120 <FEL>
                                                                     R; Izui, S. submitted to the EMBL Data Library,
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A; Residues: 1-120 < FE2>
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                                                   A; Reference number:
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Matches 98
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                                                                                                                                                                                                                                            VDDFKGRFVFSLETSASAAYLQINNLKNEDTATYFCARRGFYAMDYWGQGTSVTVSS 136
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nilarity 81.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                     S32185
                                                                                                                                                                                                                                                                                                                                                                                  41.1%; Score 517; DB 2;
80.3%; Pred. No. 2.9e-30;
ative 12; Mismatches 9
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Pred. No. 1.4
                                                                       February
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NCBIP:109961)
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N;Contains: Ig kappa chair provided Rouse Mouse)
C;Species: Mus musculus (house mouse)
C;Datc: 24-Apr-1984 #sequence revision 24-Apr-1984 #text_change 09-Jul-2004
C;Accession: A93211; B93211; A93815; A94239; A01922; A01923
C;Accession: A93211; B93211; A93815; A94239; A01922; A01923
R;Seidman, J.G.; Max, B.E.; Leder, P.
Nature 280, 370-375, 1979
A;Title: A kappa-immunoglobulin gene is formed by site-specific recombinati
A;Reference number: A93211; MUID:79221900; PMID:111146
A.Accession: A93211
A;Accession: A93211
A;Accession: A93211
A;Accession: A93211
A;Molecule type: DNA
A;Residues: 1-130 <PC41>
A;Cross references: UNIPROT:P01639
A;Cross references: UNIPROT:P01639
A;Accession: B93211
A;Accession: B93211
A;Accession: B93211
A;Molecule type: DNA
A;Residues: 1-117 <VK41>
A;Cross-references: GB:V00804; GB:J00566; NID:g52127; PIDN:CAA24186.1; PID:g575660
A;Note: the sequences were determined from the differentiated gene MOPC 41 and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig heavy chain V and J regions, monoclonal antibody SCET.M8.1 - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 20-Jun-2000 C;Accession: A53285 R;Sawada, J; Mizusawa, S.; Terao, T.; Naito, M.; Kurosawa, Y. Mizusawa, S.; Terao, T.; Naito, M.; Kurosawa, Y. Mol. Immunol. 28, 1063-1072, 1991 A;Title: Molecular characterization of monoclonal anti-steroid antibodies: primar and their pH-reactivity profiles.
A;Reference number: A53285; MUID:92017897; PMID:1922102 A;Accession: A53285; MUID:92017897; PMID:1922102 A;Ascetus: preliminary A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: DNA; protein
A;Residues: 1-119 <SAW>
A;Residues: 1-119 <SAW>
A;Cross-references: GB:D12736; NID:g220595; PIDN:BAA02228.1; PID:g220596
A;Cross-references: GB:D12736; NID:g220595; NCBIN:63271, NCBIP:63299)
A;Note: sequence extracted from NCBI backbone (NCBIN:63271, NCBIP:63299)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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81.2%;
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Pred. No. 7.8e-30;
2; Mismatches 9
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Pred. No. 5.5e-30;
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C;Superfamily: immunoglobulin V region; imm C;Keywords: heterotetramer; immunoglobulin F;12-95/Domain: immunoglobulin homology <IV
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C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kapper complex) immunoglobulin heterotetramer subunit consists of two identical light (kapper) hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into last; Superfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: alternative initiators, heterotetramer; immunoglobulin
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A;Title: Mechanism of antibody synthesis: size differences between A;Reference number: A94239; MUID:67056897; PMID:4162931
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                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-118 <WEI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Weissenhorn, W.; Riethmueller, G.; Weiss, E. submitted to the EMBL Data Library, March 1992 A;Description: Structural characterization of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig heavy chain V region (M-T406) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change
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F;3-22/Domain: signal sequence #status experimental <SIG2>
F;3-23/Domain: signal sequence #status experimental <SIG2>
F;23-130/Product: Ig kappa chain V region (MOPC 41) #status experimental <M41>
F;38-112/Domain: immunoglobulin homology <IMM>
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A;Residues: 23-49,'B',51-53,'LSB',57-58,'ZZ',61-62,'BZ',65-76,'B',78-108,110-130
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Proc. Natl. Acad. Sci. U.S.A. 74, 716-720, 1977
Prot. Natl. Acad. Sci. U.S.A. 74, 716-720, 1977
A;Tille: Anino acid sequence of the NH-2-terminal extra piece
A;Reference number: A93815; MUID:77148916; PMID:403522
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Best Local S
Matches 95
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FKGRFAFSLETSASAAFLLINNLKNEDTATYFCTRAFYDYDGAWFPYWGQGTLVTVSA 118
                                               FKERFAFSLETSATTAHLQINNLRNEDSATYFCARRF-----GFAYWGQGTLVSVSA 115
                                                                                                   LVQSGPELKTPGETVKISCKASGYTFTNYGMNMMEQAPGKGLKMMGWINTYTGEPTYADD
                                                                                                                                                        LVQSGPELKKPGETVKISCKASGYTFANYGMNWMKQAPGKGLKWMGWINTYTGQSTYADD 63
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93.5%;
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                                                                                                                                                                                                            Score 507; DB 2;
Pred. No. 1.3e-29;
6; Mismatches 11
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Pred. No. 1.2e-29;
2; Mismatches 5;
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R;Jannot, C.B.; Hynes, N.E.
Biochem. Biophys. Res. Commun. 230, 242-246, 1997
A;Title: Characterization of scFv-421, a single-chain antibody
A;Reference number: JC5322; MUID:97168950; PMID:9016757
                                                                                 p53 specific single-chain antibody Pab421 - C;Species: Homo sapiens (man) C;Date: 15-May-1997 #sequence_revision 15-Ma C;Datession: JC5322
                                                                                                                                                                             JC5322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-118 <FIS>
A;Residues: 1-118 <FIS>
A;Cross-references: EMBL:X74589; NID:g402599; PIDN:CAA52666.1; PID:g1333981
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;15-98/Domain: immunoglobulin homology <IMM>
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S37204

Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change
C;Accession: S37204
R;Fischer, R; Voss, A.; Hunziker, W.; Stierhof, Y.D.; Kreuzaler
submitted to the EMBL Data Library, August 1993
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A;Description: Production and cloning of TMV-specific monoclonal antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Status: preliminary
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C;Superfamily: immunoglobulin V region; immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin P;10-93/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: mRNA
A; Residues: 1-115 <WEI>
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C;Species: Mus musculus (house mouse)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change
C;Accession: S19968
R;Weissenhorn, W; Riethmueller, G.; Weiss, E.M.; Rieber, E.P.
submitted to the EMBL Data Library, March 1992
A;Description: Structural characterization of CD4 mAb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: preliminary
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Pred. No. 1.6e
8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   Score 503.5; DB 2
Pred. No. 2.2e-29;
.2; Mismatches 7
                                                                                                    15-May-1997
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A;Accession: JC5322
A;Mecule type: mtNA
A;Residues: 1-233 <JAN>
A;Experimental source: hy
C;Comment: This protein s
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F;98-106/Region:
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R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, J. Exp. Med. 171, 265-297, 1990

A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig kappa chain V region (anti-DNA, DP7VK) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Dspecies: 16-Sep-1992 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                      F;24-34/Region: complementarity-determining F;35-49/Region: framework 2
                                                                                                                                                                                                                                                                                                                                                                        C; Superfamily: immunoglobulin V region; immunoglobulin C; Keywords: heterotetramer; immunoglobulin
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;Comment: This protein specifically binds the tumor suppressor protein p53. It restores
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61 RFSGSRSGSDYSLTISSLESEDFVDYYCLQYASSPWTFGGGTKLEI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       103;
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                                                                     DIQMTQSPSSLSASLGERVSLTCRASQDIGSSLNWLQQEPDGTIKRLIYATSSLDSGVPK
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framework 3
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43.3%; Pred. No. 7.3e-29;
tive 46; Mismatches 79
                                                                                                                                                  2;
                                                                                                                                                                  Score 498; DB 2;
Pred. No. 4.9e-29;
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Ig kappa chain V region (anti-DNA, DP11VK) - mouse c;Species: Mus musculus (house mouse) C;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 C;Accession: PL0259
C;Accession: PL0259
C;Shlomchik, M.; MasCelli, M.; Shan, H.; Radic, M.Z J. Exp. Med. 171, 265-297, 1990
A;Title: Anti-DNA antibodies from autoimmune mice a

autoimmune mice arise by clonal expansion and somatic

Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein,

#text_change 21-Jan-2000

(fragment)

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Ig heavy chain V region - mouse
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C;Accession: 826325
R;Stark, S.E.; Caton, A.J.
J. Exp. Med. 174, 613-624, 1991
A;Title: Antibodies that are specific for a single amino acid interchange in a protein A;Reference number: $26309; MUID:91341421; PMID:1908510
A;Accession: $26325
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A,Molecule type: mRNA
A,Residues: 1-109 <STA>
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F;16-90/Domain: immunoglobulin homology <IMM>
F;24-34/Region: complementarity-determining 1
F;35-49/Region: framework 2
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A;Residues: 1-106 <SHL>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
Search completed: April 4,
Job time: 89.7284 secs
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P;57-88/Region: framework 3
P;89-97/Region: complementarity-determining 3
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Matches
                                                                                                                                                                                                                                                                                               Query Match 39.5%; Score 497.5; DB 2; Best Local Similarity 85.3%; Pred. No. 5.5e-29; Matches 93; Conservative 7; Mismatches 4;
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Local Similarity 93.4%;
hes 99; Conservative
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Pred. No. 4.9e-29;
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2: uniprot_trembl:*
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Q65ZC8
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Q65ZL2
Q96KB05
Q65ZC7
Q99LA6
KV5G_MOUSE
Q925S9
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Q65zc9 homo sapien
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Q7tom2 mus
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Q65212 mus
Q68205 mus
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Q95189 mus
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Matches 161;
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ESQUENCE FROM N.A.

MEDLINE=98170165; PubMed=9509426;

Chung J.H., Choi S.J., Kim H.J., Kim I.J., Choi I.H., Lee S.D.

A Yi K.S., Suh P.G., Ryu S.H., Chung H.K.;

A Yi K.S., Suh P.G., Ryu S.H., Chung H.K.;

"Cloning and characterization of cDNAs encoding VH and VL of immonocional anti-CEA antibody (CEA 79) cross-reactive with NCA generation of a single-chain Fv molecule (scFv).";

Mol. Cells 7:816-819(1997).

R EMBL; U88067; AAB48044.1; -.

R EMBL; U88067; AAB48044.1; -.

R PIR; S19965; S19965.

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R PIR; S19968; S19968.

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R PIR; S19968; S19968.

PIR; S19968; S19967.

PIR HSSP; PO1607; ABMPA.

PIR HSSP; P
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Q921A6;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-CEA 79 single chain Fv (Fragment).
Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                 LEIKRA 242
                                                                                                                                             ISSSSGSSSSGSSSSGSDIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGT
                                                                                                                                                                                                                                                                                  QVKLQQSGPELKKPGETVKISCKASGYTFTDYGMNWVKQAPGKGLKWMGWINTYTGEPTY
                                         PRSAHTLHIYIQPGIPSRFSGSGSGRDYSFSISNLEPEDIATYYCLHYDNL-HTFGGGTK 235
                                                                        IKRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTN
                                                                                                                      -SGGGSGGGGSGGGSDIELTQSPSSLSASLGGKVTITCKASQDINKYIAWYQHKPGKG
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                                                                                                                                                                                                                                                                                                                                                                                                                       241
26086 MW; 0276887248E9C771 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65.8%; Score 827.5; DB 2
65.4%; Pred. No. 3.9e-58;
tive 27; Mismatches 49
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KOUSE
Q6GWI
Q7Z3Y4
Q7Z3Y4
Q6GWX8
Q6GWX8
Q6GWX0
Q91WR1
Q6GWX0
Q91WR1
Q91W71
KV1G_HUMAN
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Q6gmw1
Q7z3y4
Q6gmx8
Q6gmx0
Q6gnx0
Q6mv1j1
Q8v1j1
Q8v1j1
Q8v1j1
Q91wt1
Q91wt1
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Q91s0362
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homo sapien
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OCCOSSION DATE
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                                                                               Q65ZC9 PREMERTED 28, Cr
Q65ZC9; CTEMBLrel 28, Cr
25-OCT-2004 (TrEMBLrel 28, Lr
T 25-OCT-2004 (TrEMBLrel 28, Lr
T 25-OCT-2004 (TrEMBLrel 28, Lr
Cingle-chain Fv (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 137
         Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON_TER
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SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 2.
PROSITE; PS50835; IG_LIKE;
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"Complement recruitment using bispecific
Nat. Biotechnol. 15:629-631(1997)
EMBL; Y13057; CAA73500.1;
InterPro; IPR00359; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR00356; Ig_v.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-OCT-2004 (TrEMBLrel. 28, 25-OCT-2004 (TrEMBLrel. 28, 25-OCT-2004 (TrEMBLrel. 28, Single-chain Fv (Fragment).
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137; Conserv
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|ELKRA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    58.6%; Score 737.5; DB 2; 55.5%; Pred. No. 6.1e-51; tive 47; Mismatches 54;
                                                                                                                              Last sequence update)
Last annotation update)
                                                                                                                                                                              Created)
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Last annotation update)
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         Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Best Local S
Matches 135
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"Complement recruitment using bispecific di.
Nat. Biotechnol 15:629-631(1997).
EMBL; Y13056; CAA73499.1; -.
Interpro; IPR003599; Ig. -.
Interpro; IPR0077110; Ig-like.
Interpro; IPR007510; Ig-v.
Pfam; PF00047; ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 2.
SMART; SM00406; IGv; 2.
PROSITER; P850835; IG_LIKE; 2.
NON_TER
                                                                                                                                                                                                                                     Q925S1
Q925S1;
Q1-DEC-2001
Q1-DEC-2001
Q1-OCT-2003
                   STRAIN-BALB/c;
Cui D., Zeng G., Yan X., Li
"Cloning of mouse genes rela
of the irradiated mice by tr
                                                                                "Mechanism of exogenous nucleic acids and their precursors improving the repair of intestinal epithelium after irradiation in mice."; world J. Gastroenterol. 6:709-717(2000).
                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
            the same strain."
                                                               SEQUENCE FROM N.A.
                                                                                                                            PubMed=11819679;
Cui_D., Zeng G.,
                                                                                                                                                  STRAIN=BALB/c;
                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                NCBI_TaxID=10090;
                                                                                                                                                                                                                 Mus musculus (Mouse)
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NON TER
SEQUENCE
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SEQUENCE FROM N.A.
STRAIN-C1q/7;
STRAIN-C1q/7;
MEDLINE-97362799; PubMed-9219263;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            240 AA;
                                                                                                                                                                                                                                                                               PRELIMINARY;
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Rodentia;
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25569 MW;
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55.6%; Pred. No. 3.4e-47;
tive 40; Mismatches 63
 Relat.
                  i., Li X., Su C.;
is related to repairing
by treatment with the
                                                                                                                            Wang
                                                                                                                                                                                                                                                           Created)
                                                                                                                                                                                                                                    Last sequence update)
Stud. Phys.
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                                                                                                                                                                                          Craniata; Vert
Sciurognathi;
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hi; Muridae; Murinae; Mus
Chem.
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                   of intestinal e intestinal RNA
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A of mice o
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ID Q7TQ
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Best Local &
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HSSP; P01751; 1A6W.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 2
PROSITE; PS50835; IG_LIKE; 2.
NON_TER 1 1
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Q7TQM2;

O1-OCT-2003 (TrEMBLrel. 25,

O1-OCT-2003 (TrEMBLrel. 25,

O1-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE-22853226; PubMed=12860977; DOI=10.1074/jbc.M306877200; Peter J.C., Eftekhari P., Billiald P., Wallukat G., Hoebeke J.; "scrv single chain antibody variable fragment as inverse agonist the beta-2 adrenergic receptor."; the beta-2 adrenergic receptor."; J. Biol. Chem. 278:36740-36747(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ScFv 6H8 protein (Fragment).
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HSSP; P01665; 1QNZ.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1
                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN-Balb/C
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    118
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ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCAR-RFGFAYWGQCTLVSVSASISS 119
                                                                                                                                                                                                              EIQLVQSGPELKKPGETVKISCKASGYTFANYGMMMKQAPGKGLKMMKWINTYTGQSTY
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                                                                                         DEKFKNKGILTVDTSSSTAYMHLSSLASEDSAVYYCARGGRGLDVWGAGTTLTVS---SG
                                                                                                                                                                                      QVQLQQSGSELVRPGASVKLSCKASGYTFTTYWMHWVKQRHGQGLEWIGNIYPGSGITNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SGGGGSGGGGSGSGSDÍVLTÓSPASLAVSLGQRATISCRÁSESVDNIG: (SFMNWFQQKP 179
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218 AA;
                                                                                                                                                                                                                                                                                                                                                                            243 AA; 25976 MW;
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58.9%; Pred. No. 6.9e-45;
tive 32; Mismatches 48
                                                                                                                                                                                                                                                                             50.6%; Score 636.5; DB 2; 52.3%; Pred. No. 7.1e-43; tive 43; Mismatches 67;
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                            BEFFF64D2DCF4F76 CRC64;
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  RESULT
Q65ZL2
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Q9QYF0;
Q9QYF0;
Q1-MAY-2000 (TrEMBLrel. 13, C
01-MAY-2000 (TrEMBLrel. 13, L
01-OCT-2003 (TrEMBLrel. 25, L
CN 8 single chain antibody.
Name=CN 8 scFv;
synthetic construct.
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 2.
PROSITE; PS50835; IG_LIKE; 2.
SEQUENCE 298 AA; 31867 MW;
Q65ZL2;
Q65ZL2;
25-OCT-2004
25-OCT-2004
25-OCT-2004
FV/M4.
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PIR; S19112; S19112.
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EMBL; AB036341; BAA88633.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   recognizing a cell polarity by using method.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shinohara N., Demura T., Fukuda H.; "Isolation of a vascular cell wall-specific monoclonal antibody
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                                                                                                                                                                                                                                                                                                                                                                                         KRLIYATSSLDSGVPKRFSGSRSGSDYSLTISSLESEDFVVYYCLQYAIFPYTFGGGTNL
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SGGGSGGGGGGGDIELTQSPASLSASVGETVTITCRASGNIHNYLAWYQQKQGKSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SSSSGSSSSGSSSIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTI 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TPSLKDKFIISRDNAKNTLYLQMSKVRSEDTALYYCARASYYGHSAYWGQGTTVTVS---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDEKERFAESLETSATTAHLQINNLRNEDSATYFCARR--FGF-AYWGQGTLVSVSASI 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QVKLQQSGGGLVKPGGSLKLSCAASGSDFSRYWMSWVRQAPGKGLEWIGEINPDSSTINY
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                                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                            281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50.6%; Score 636; DB 2; Length 2:
51.4%; Pred. No. 1e-42;
tive 42; Mismatches 71; Indels
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Last annotation update)
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(TrEMBLrel. 28, (TrEMBLrel. 28, (TrEMBLrel. 28,

Last sequence update)
Last annotation update)

Created)

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RESULT
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Best Local S
Matches 122
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Q6KB05;
05-JUL-2004 (TrEMBLrel. 27, Cre
05-JUL-2004 (TrEMBLrel. 27, Las
05-JUL-2004 (TrEMBLrel. 27, Las
SCFV B8E5 protein (Fragment).
Name=scFv B8E5;
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InterPro; IPR007110; Ig-1ike.
InterPro; IPR003597; Ig-c1.
InterPro; IPR003598; Ig-c2.
InterPro; IPR003598; Ig-MHC.
InterPro; IPR003596; Ig-MHC.
InterPro; IPR003596; Ig-W.
Pfam; PF07554; C1-set; 2.
Pfam; PF00047; ig-4.
SMART; SM00409; IG-3.
SMART; SM00409; IG-3.
SMART; SM00409; IG-2; 2.
SMART; SM00408; IG-2; 2.
SMART; SM00408; IG-2; 2.
PROSITE; PS50835; IG-LIKE; 4.
PROSITE; PS50835; IG-MIC; UNIXNO
SEQUENCE 487 AA; 53578 MW;
                                 STRAIN=Balb/c;
Peter J.C., Wallukat G., Tugler J., Maurice D.
Briand J.P., Hoebeke J.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ
EMBL, AJ746180; CAG34081.1; -.
HSSP; PO1837; NRCR.
                                                                                                                                                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
       InterPro;
InterPro;
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Mus sp.
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                               NCBI_TaxID=10090;
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"A genetically engineered single-gene-encoded antibody secreted from myeloma cells.";
Hum. Antibodies Hybridomas 6:161-166(1995).
EMBL; S82493, AAB37424-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=96272580;
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  IPR003599;
IPR007110;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VYAATNLADGVPSRFSGSGSGTQYSLKINSLQSEDFGSYYCQHFWGTPYTFGGGTRLEIK
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Ig.
Ig-like
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50.8%; Pred. No. 6.1e-42;
tive 46; Mismatches 67
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RESULT 9
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AC Q65
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MEDLINE=52020904; PubMed=1924323;

MEDLINE=52020904; Pai L.H., FitzGerald D.J., Willingham M.,

Brinkmann U., Pai L.H., FitzGerald D.J., Willingham M.,

"B3(Fv)-FB38KDEL, a single-chain immunotoxin that cause

regression of a human carcinoma in mice.";

regression of a human carcinoma in mice.";

regression of a human carcinoma in mice.";

REMBL; S57990; AAB19971.2;

REMBL; S57990; AAB19971.2;

RINGEPTCO; IPR003599; IG.

RINGEPTCO; IPR003599; IG.

RINGEPTCO; IPR003596; IG_v.

Pfam; PF00047; IG; 2.

SMART; SM00409; IG; 2.

SMART; SM00409; IG; 2.

SMART; SM00409; IG; 2.

RONGITE; PS50835; IG LIKE; 2.

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248

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RONGITE; PS50835; IG LIKE; 2.
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Best Local Similarity
Matches 121; Conserv
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Q65ZQ7;
25-OCT-2004
25-OCT-2004
25-OCT-2004
                                                                                                                                                                                                                                                                      SEQUENCE
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Mammalia; Eutheria;
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25-OCT-2004 (TrEMBLrel. 28,
25-OCT-2004 (TrEMBLrel. 28,
83(FV)-PE40 (Fragment).
Name=B3(FV)-PE40;
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SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 2.
PROSITE; P850835; IG_LIKE;
NON_TER 1
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                                                                                                                                                                       120;
61 ADDFKERFAFSLETSATTAHLQINNLRNEDSATYFCARRFG----FAYWGQGTLVSVSAS
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                                                                DVKLVESGGGLVQPGGSLKLSCATSGFTFSDYYMYWVRQTPEKRLEWVAYISNDDSSAAY
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                                                                                                                                                                                                                                                                 248 AA;
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Rodentia;
                                                                                                                                                                                           46.9%;
                                                                                                                                                                                                                                                                 26634 MW;
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                                                                                                                                                                  Score 590; DB
Pred. No. 3.8e
45; Mismatches
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1.8e-39;
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RESULT 10
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Brownstein M.J., Uddin T.B., Toshiyuki S., Caraninci P., Prange C.,

A Roman S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Phbey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Phhey J., Helton E., Ketteman M., Madan A., Schein J.S.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Query Match
Best Local Similarity
                                                                                                                                                        InterPro; IPR007110; Ig-11ke.
InterPro; IPR003597; Ig C1.
InterPro; IPR003597; Ig_MHC.
InterPro; IPR003906; Ig_MHC.
Pfam; PF07654; C1-set; Z.
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Q99LA6;
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MEDLINE-22388257; PubMed-12477932; DOI-10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
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01-JUN-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                            PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
Hypothetical protein.
SEQUENCE 484 AA; 52567 MW; BEAEA
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Mammalia; Eutheria; Rodentia;
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SMART; SM00406; IGv; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jones S.J., Marra M.A.; "Generation and initial analysis of more and mouse cDNA sequences.";
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MEDLINE=67056897; PubMed=4162931; Gray W.R., Dreyer W.J., Hood L.; "Mechanism of antibody synthesis:
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MEDLINE=79221900; PubMed=111146;
Seidman J.G., Max E.E., Leder P.;
Seidman J.G., Max E.E., Leder P.;
"A kappa-immunoglobulin gene is formed without further somatic mutation.";
Nature 280:370-375(1979).
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P01639; P01640;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
PROSITE; PS50835; IG_LIKE; 1.
Bence-Jones protein; Direct protein sequencing;
Immunoglobulin V region; Signal.
                                                                        InterPro; IPR007110; Ig-like
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Burstein Y., Schechter I.; "Amino acid sequence of the NH2-terminal extra piece segments of precursors of mouse immunoglobulin lambdal-type and kappa-type 1:
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MEDLINE=77148916; PubMed=403522;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                   PIR; A93211; KVMSM4.
                                                                                                                                                                                                                                                                                                     Science 155:465-467(1967).
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Matches 98
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Best Local Sim
Matches 101;
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Q925S9;
01-DEC-2001
01-DEC-2001
01-OCT-2003
Q683Y7;
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DISULFID
VARIANT
NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=99306687; PubMed=10380019;
Tripathi P.K., Qin H., Bhattacharya-Chatterjee M., C
Foon K.A., Chatterjee S.K.;
"Construction and characterization of a chimeric fus
consisting of an anti-idiotype antibody minicking a
associated antigen and the cytokine GM-CSF.";
Hybridoma 18:193-202(1999).
BMBL; AF124721; AAK55120.1; -.
BMSL; AF124721; AKS5120.1; -.
BMSL; SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequer
01-OCT-2003 (TrEMBLrel. 25, Last annote
Immunoglobulin light chain (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata;
Mammalia; Butheria; Rodentia; Sciurogna
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NON TER 127 127
SEQUENCE 127 AA; 13794 MW;
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                                                                                                                                                                                                                                                                                    98;
                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                          DIQMTQSPSSLSASLGERVSLTCRASQDIGNSLTWLQQEPDGTIKRLIYATSSLDSGVPK
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                                                                                                                                                                                                 DIQMTQSPSSLSASLGQRVSLTCRASQDIGINLHWLQQEPDGTIKRLIYATSSLGSGVPK
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                                                                                                                      RFSGSRSGSDYSLTISSLESEDFVAYYCLQYASSPYTFGGGTKLEIK 127
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                    PRELIMINARY;
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Pred. No. 6e-33;
2; Mismatches 5
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Complementarity-determining-2.
Framework-3.
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                    PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
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                    116
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..1e-31;
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DT 25-QC
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Best Local
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                          plasmon resonance analysis.";
J. Immunol. Methods 193:177-187(1996).
EMBL; S82492; AAB37434.2; -.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig-like.
InterPro; IPR00359; Ig-v.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
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InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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                                                                                                                                                                                 SEQUENCE FROM N.A.

MEDLINE=96269537; PubMed=8699031; DOI=10.1016/0022-1759(96)00043-9;

Noel D., Bernardi T., Navarro-Teulon I., Marin M., Martinetto J.P.,

Ducancel F., Mani J.C., Pau B., Piechaczyk M., Biard-Piechaczyk M.;

"Analysis of the individual contributions of immunoglobulin heavy and light chains to the binding of antigen using cell transfection and
                                                                                                                                                                                                                                                                                                                                                                                                              25-OCT-2004 (TrEMBLrel.
25-OCT-2004 (TrEMBLrel.
25-OCT-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                              Name=Tg10H;
                                                                                                                                                                                                                                                                                                                                                                                               Tg10H (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q65ZL3;
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Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ784033; CAH04483.1; -.
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STRAIN=Balb/c;
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-OCT-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Immungolobulin heavy chain variable region (Fragment).
                 NON_TER
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 140 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                              Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12863 MW;
   15302 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                               228
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Sciurognathi; Muridae;
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Sciurognathi; Muridae;
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   7BFE237F3D965F43 CRC64;
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; Murinae; Mus.
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Q9JL79
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Search completed: April 4, Job time: 412.693 secs
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STRAIN-A.CA;

MEDLINE-20448942; PubMed=10992488;

MEDLINE-20448942; PubMed=10992488;

DOI=10.1128/IAI.68.10.5803-5808.2000;

MAIkiel S., Liao L., Cunningham M.W., Diamond B.;

"T-Cell-dependent antibody response to the dominant epitope of streptococcal polysaccharide, N-acetyl-glucosamine, is cross-reactive with cardiac myosin.";

Infect. Immun. 68:5803-5808(2000).

EMBL; AP206027; AAF89325.1; -.

HSSP; P01751; 1NOB.
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Q9JL79;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Anti-myosin immunoglobulin heavy chain variable region
(Fragment).
(Fragment).
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InterPro; IPR003596; Ig_v.
BMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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New single chain Fv polypeptide binding to C-erbB-2 tumour antigen - imaging or treating breast or ovarian cancer etc. Example; Page 70; 87pp; English. N-PSDB; Huston JS, 1993-272889/34. DB; AAQ46087. Houston LL, Ring DB, Oppermann ĸ

for

c-erbB-2 refers to a protein antigen expressed on the surface of tumour cells. such as breast and ovarian tumour cells, which is an approx.

200,000 mol. wt. acidic glycoprotein having an isoelelectric pt. of about 5.3 (see AAQ46083, AAR39568). A single chain Fv (sFv) is a covalently linked VH-VL heterodimer which is expressed from a gene fusion including VH- and VL- encoding genes connected by a peptide-encoding linker. Such linker sequences are set forth in AA residues 116-135 in AAR39569, which includes part of the 16 AA linker sequences in AAR39572. Using AAQ46084 for the 520C9 monoclonal antibody, a single chain polypeptide can be produced having a binding affinity for a c-erbB-2 related antigen.

(Updated on 25-MAR-2003 to correct PN field.)

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RESULT 2
AAY90828
AAY90828
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                                                                                                                                                                                                                                  The present invention describes a monoclonal antibody (MAb) (I) that binds to a human breast cancer antigen that is also bound by MAb 454C11 and 52CC9 (produced hybridoma ATCC HB8484 and HB8696, respectively). Also described is a hybridoma that produces (I). (I) is useful in specific binding assays, affinity purification, drug or toxin targeting, imaging, and genetic or immunological therapeutics for various cancers. The present sequence represents a linker amino acid sequence which is used in the exemplification of the present invention
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21-MAR-1986;
08-MAY-1988;
                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Col 77; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monoclonal antibody useful for affinity
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                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 4
AAU04949
ID AAUC
XX
AC AAUC
XX

AAU04949 standard;

peptide; 15

AA

AAU04949;

밁 S

Matches Query Match Best Local :

. Similarity 15; Conserv

100.0%; llarity 100.0%; Conservative 0,

0

Score 66; DE Pred. No. 0.0); Mismatches

DB 4; 0.012;

Length 15;

0

0,

Gaps

0

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RESULT 3
AAU08688
ID AAU08608
ID AAU0
XX AAU0
AX AAU0
XX Angi

                                                   The invention relates to a conjugate comprising a molecule that exerts a biocidal or cytotoxic effect on target cells in lesions of pathological angiogenesis and an antibody (L19) directed against an extracellular matrix component (the ED-B domain of fibronectin) which is present in the lesions. Examples of biocidal or cytotoxic molecules include fibronectin-gamma), Tumour Necrosis Factor alpha (II--12), Interferon-gamma (IFN-these proteins may interact with a membrane-bound receptor on the target cell or perturb the electrochemical potential of the cell membrane. The restable disorders include tumours, rheumatoid arthritis, diabetic retinopathy, age related muscular degeneration and angiomas. This sequence represents an amino acid linker encoded by a PCR primer used in the construction of an L19-II-2 fusion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zardi L,
Halin C;
   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel conjugate for treating lesions, comprises a specific binding specific for extra-cellular matrix component present in lesions, are molecule that exerts biocidal/cytotoxic effect on target cells in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Angiogenesis; extracellular matrix; ED-B domain; fibronectin; lesion; IL-2; doxorubicin; interleukin-12; Interferon-gamma; IL-12; IFN-gamma; Tumour Necrosis Factor alpha; TNF alpha; tissue factor protein; angioma; electrochemical potential; tumour; cancer; rheumatoid arthritis; diabetic retinopathy; age related muscular degeneration; cytostatic; antirheumatic; antiarthritic; antidiabetic; ophthalmological.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-FEB-2000;
21-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-FEB-2001; 2001WO-IB000382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lesions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PHIL-) PHILOGEN SRL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200162298-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody L19-Fibronectin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU08688;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU08688 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-596713/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Page 30; 88pp;
   15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neri
   AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0184767P.
2000US-0257192P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Carnemolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (IL-2) fusion protein amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Å
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nilsson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Borsi
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24-OCT-2001

(first entry)

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ABW00718
ID ABW0
XX ABW0
XX ABW0
XX Link
XX Cink
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XX Cell
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                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                     The sequence represents a linker peptide used to make a humanised pl85 (520C9) single chain antibody/human interleukin-2 (IL-2) fusi protein. The fusion protein (or immunoconjugate) is used to inhibit overexpression of pl85 e.g. human ademocarcinomas and malignant an benign tumours of the breast, renal system, salivary gland, gastrointestinal tract or gastric tumours
          06-FEB-1992;
07-OCT-1993;
                                                                    14-NOV-2002
                                                                                          US2002168375-A1
                                                                                                                 Unidentified
                                                                                                                                   Cell proliferation;
                                                                                                                                                            Linker #2,
                                                                                                                                                                                   15-JAN-2004
                                                                                                                                                                                                         ABW00718;
                                                                                                                                                                                                                               ABW00718
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 16; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel immunoconjugate useful for inhibiting tumor cell growth in vivo comprises a humanized anti-p185 antibody linked to an Interleukin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-451904/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Austin R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-JAN-2000; 2000US-0177258P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HAMI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHIR )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Humanised;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Humanised anti-p185 antibody/IL-2
                                                                                                                                                                                                                                                                                                                                     15;
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                                                                                                                                                                                                                                                                                                              <u>بــ</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHIRON CORP.
HAMILTON CIVIC HOSPITALS
HAMILTON REGIONAL CANCER
                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                       SSSSGSSSSGSSSG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody; p185; 520C9H; interleukin-2; IL-2; mour; adenocarcinoma; fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kwok CS,
                                                                                                                                                           peptide
                                                                                                                                                                                                                                                                                                                                     Conservative
                                            2001US-00887853.
                                                                                                                                                                                   (first entry)
92US-00831967.
93US-00133804.
95US-00462641.
                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                        used in the invention.
                                                                                                                                                                                                                              peptide;
                                                                                                                                    cytotoxic agent; drug targetting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ring DB;
                                                                                                                                                                                                                                                                                       15
                                                                                                                                                                                                                               15
                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                               Score 66;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RES DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fusion
                                                                                                                                                                                                                                                                                                                                                 0.012;
                                                                                                                                                                                                                                                                                                                                                          DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein,
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                                                                                                                                                                                                                                                                                                                                                          Length 15;
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                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunoconjugate;
                                                                                                                                                                                                                                                                                                                                   <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                    inhibit the
                                                                                                                                                                                                                                                                                                                                                                                                                                                              fusion
                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                             and/or
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RRESULT 6
ADN66232
ID ADN66
XX ADN66
XX ADN66
XX Iink
XX Iink
XX Iink
XX Ion
DE Co-J
PD 20-A
PD 20-A
PF 02-J
XX WPI;
DE E
PT Comp
PT Dete
PT Comp
PT Dete
PT Comp
PT Dete
PT Comp
PT Toete
PT Dete
PT Comp
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a binding protein formulation for targetting epitope on an antigen expressed in mammal. The formulation comprises c dimeric biosynthetic construct having conformation permitting binding antigen by binding site of each polypeptide chain. The invention is useful for imaging a preselected antigen in a mammal expressing the antigen. It is also useful for targetting drugs that inhibits cell proliferation and cytotoxic agents that kill cells. The present sequen is a linker peptide used in the invention
                                                                                                                                                                                                     Determining of distance between binding sites on target molecule comprises contacting target molecule with self-assembled monolays identifying controlled geometry with lowest density.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel binding protein formulation for targeting epitope on antigen in mammal, comprising dimeric blosynthetic construct having conformation permitting binding of antigen by binding site of each polypeptide chain.
                        binding sites on a target molecule, involving contacting the tarmolecule with a self-assembled monolayer comprising binding mois immobilised with respect to each other in controlled geometries.
                                                                                                                                                        Disclosure; SEQ ID NO 18; 37pp; English.
                                                                                                                                                                                                                                                                                                              WPI; 2004-386879/36.
                                                                                                                                                                                                                                                                                                                                                                Bamdad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6723517-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Linker peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADN60232;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADN60232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15
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                                                                                                     The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linker; self-assembled monolayer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-APR-2000; 2000US-00558741.
                                                                                                                                                                                                                                                                                                                                                                                                                (MINE-) MINERVA BIOTECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     $$$$G$$$$G$$$$G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SSSSSSSSSSSSG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
larity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 2; 30pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                  relates to a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0087766P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9905-00324258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 66; DB 7;
Pred. No. 0.012;
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                                                                                                                                                                                                                                                                                                                                                                                                                   CORP
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                                                                                                     for determination
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7;
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H
                                                                                                        distance between
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                                                        moieties
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                                                                                                                                                                                                                                        and
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identifying the controlled geometry with

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RESULT 7
ADR38601
ID ADR3
XX ADR3
AC A
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The invention describes an isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a specific clone where (I) binds to and neutralises botulinum neurotoxin type A (BONT/A). An isolated antibody (I) that specifically binds to an epitope specifically bound by an antibody expressed by a clone chosen from clone specifically bound by an antibody expressed by a clone chosen from clone S25, C25, C39, IC6, 3D12, B4, IF3, huC25, Ar1, Ar2, WR1(V), WR1(T), 3-1, 3-8, 3-10 and ING1, where (I) binds to and neutralizes botulinum neurotoxin type A (BONT/A). Also described are: a polypeptide (II)
                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated antibody that neutralizes botulinum neurotoxin type A, useful for diagnosing botulism or for treating pathologies associated
                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 3; 110pp; English.
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01-AUG-2002; 2002US-0400721P.
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albumin (EPOa-hSA) fusion proteins having at least one amino acid residue of the EPOa moiety altered such that a site, which serves as a site for glycosylation in EPO does not serve same purpose in the EPOa. The present sequence represents a specifically claimed peptide linker for use in the construction of an EPOa-hSA fusion protein. The EPOa-hSA fusion protein

The present invention describes erythropoietin analogue-human

(EPOa-hSA)

An erythropoietin analog-human serum albumin fusion protein, e.g. treatment of HIV infection.

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06-OCT-2000;
29-DEC-2000;
                         The present invention relates to a nucleic acid comprising a sequence encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, seppsis, cirrhosis, dermatitis, psoriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveitis and coeliac disease. The present sequence is a linker peptide described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; rheumatoid arthritis; asthmacil tribosis; dermatitis; psoriasis; inflammatory bowel disease; immunosuppressive; antinflammatory; antirheumatic; antiarthritic; antiasthmatic; antibacterial; dermatological; antipsoriatic; antidiabettic; ophthalmological; neuroprotective; multiple sclerosidiabetes; uveitis; coeliac disease; linker.
                                                                                                                                                                                                                           Novel nucleic acid encoding fusion protein stimulating hormone concatamer or its analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              can be used in treatment of a subject in need of EPO, e.g. for a patient suffering from anaemia associated with renal failure, chronic disease, HIV infection, blood loss or cancer. The EPOa-hSA fusion protein can be included in a nutraceutical, such as an infant feeding formula. The EPOa-hSA constructs can be used as part of a gene therapy protocol to delivery nucleic acids encoding an EPOa-hSA fusion protein
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                                                                                                                                                                                    Disclosure;
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                                                                                                                    The invention relates to a method of generating an antibody. The method is useful in generating antibodies that may be used for basic research and in the diagnosis and treatment of various diseases. The antibodies may allow for the precise identification and quantitation of a particular species of protein in a biological sample. The antibody may also be used to modulate the activity or function of a protein which is useful for treating a disease characterised by excessive or insufficient activity of the protein. The method allows for the rapid and efficient production and isolation of large amounts of protein or its portion in its native conformation. The method avoids the time-consuming, expensive and laborious tasks associated with the in vitro production and purification of proteins. The present sequence represents a peptide linker.
                                                                                                                                                                                                                                                                                                                   Generating an antibody for basic research and therapeutic purposes, comprises producing a fusion protein in a first mammal and administering the formed protein to a second mammal to generate an antibody against the
                                                                                                                                                                                                                                                                                                                                                                                             ₩u C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17
                                                                                               Sequence
                                                                                                                                                                                                                                                                                 Disclosure; Page 3; 9pp; English.
                                                                                                                                                                                                                                                                                                          protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  (WUCC/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibody generation; rapid protein production; efficient protein production; linker.
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                                                                                                                                                                                                                                                                                                                                                                                                                       (HSUP/)
                                                                                                                                                                                                                                                                                                                                                                                             Lin R,
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LIN R.
HSU P.
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                                                             Similarity
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                                               Conservative
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                                                           Score 66; DB 6;
Pred. No. 0.014;
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Pred. No. 0.014;
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                                                 Mismatches
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RESULT 11
ADB24915
ID ADB24
AC ADB24
AC
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ABU10017
        ZZXBXBXBXB
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Best Local S
Matches 15
Bladder disorder; un-methylated CpG
                                                                                Alpha-MSH concatamer related linker peptide
                                                                                                                                                  31-JUL-2003
                                                                                                                                                                                                                                                          ABU10017 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of preparing a purified fusion protein. The method is useful for preparing fusion proteins. The method allows for rapid and efficient, production and isolation of a protein encoded by any nucleic acid sequence of interest. The method permits the high-throughput screening of antibodies or other ligands directed against any protein sequence, without the need to synthesise the protein sequence in vitro. The method avoids the time-consuming, expensive and laborious tasks associated with the in vitro production and purification of proteins. The method avoids several of the difficulties associated with recombination protein production in bacteria. The present sequence represents a peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WUCC/) WU C.
(LINR/) LIN R.
(HSUP/) HSU P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 3; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preparing purified fusion proteins useful to screen for target binding molecules, by administering nucleic acid encoding the fusion protein to mammal, and isolating the fusion protein by removing a biological sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fusion protein efficient prote
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-2001; 2001US-0318474P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide linker #4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SSSSGSSSSGSSSSG
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ilarity 100.0%; !
Conservative 0;
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sin production; high-throughput screening; linker
cytostatic; antiinflammatory; immune respon sequence; alpha-MSH; melanocortin receptor;
                                                                                                                                                                                                                                                                                                                                                                                                      15
                                                                                                                                                                                                                                                                                                                                                                                                                                                            15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 66; DB 7;
Pred. No. 0.014;
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                       immune response;
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The invention describes a method of modulating an immune response in a CC mammal, comprising identifying a mammal that has or is at risk for having a bladder disorder, and administering: (a) an isolated mucleic acid (N1) CC comprising an un-methylated CpG sequence to the mammal; (b) an isolated CC nucleic acid (N2) comprising sequence encoding alpha-MSH to the mammal; CC method is useful for modulating immune response in a mammal having a CC bladder disorder, where administration of (N1) results in an amelioration of one or more symptoms of the disorder. Preferably, the method is useful for modulating immune response in a mammal having a CC diministration of (N1) results in a decrease in tumour size or activity), or for modulating immune response in a mammal having bladder cancer (where administration of (N1) results in a modulation of the immune response to a Th1 response). The method is also useful for modulating immune response to a Th1 response). The method is also constituted that is characterised by inflammation which is associated with symptoms of interstitial cystitis or associated with a disruption of the integrity of the bladder lining. This is the amino acid sequence of a linker constituted in the creation of melanocyte stimulating hormone (alphance).
     Matches
                     Query Match
Best Local
                                                                           Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 5; 17pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modulating immune responses in a mammal with a bladder disorder e.g. bladder cancer, by administering nucleic acids comprising un-methylated bladder cances, nucleic acids encoding alpha-MSH, or alpha-MSH peptides to the sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-447327/42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bladder cancer; tumour; interstitial cystitis; inflammation; alpha-MSH; melanocyte stimulating hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2002193332-A1
                                                                                                                 concatamers used in the method of treating bladder
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                     Similarity
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   Conservative
                   100.0%;
   0
                   Score 66;
Pred. No.
   Mismatches
                   DB 7;
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                                     Length 17
   Indels
                                                                                                                   disorders
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Gaps
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RESULT 13
ADQ28276
ID ADQ28
XX
AC ADQ28
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DT 09-SE
DT Pepti
XX
Pepti
XX
Gene
KW gene
KW gene
KW prost
XX
XX
SYnth
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                                      gene therapy; transgenic; fusion protein; colon cancer; breast cancer;
prostate cancer; ovarian cancer; endometrial cancer.
                                                                                                       09-SEP-2004
                                                                                                                                                        ADQ28276 standard; peptide;
                                                                             Peptide linker #4
                                                                                                                                 ADQ28276;
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Synthetic

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RESULT 14
AAB15670
ID AAB15
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                                                                     23-JAN-1999;
03-MAY-1999;
07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of making a transgenic fusion protein comprising providing a transgenic animal which includes a transgene which provides for the expression of the fusion protein, allowing the transgene to be expressed, and recovering the fusion protein, from the milk of the transgenic animal. The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of colon, breast, prostate, ovarian and endometrial cancers. The present sequence represents the amino acid sequence of a peptide linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Making a transgenic fusion protein using a transgenic animal having a transgene expressing the fusion protein, useful for the diagnosing, preventing or treating colon, breast, prostate, ovarian and endometrial
                                                                                                                                                                           21-JAN-2000; 2000WO-US001504
                                                                                                                                                                                                                            27-JUL-2000
                                                                                                                                                                                                                                                                          WO200043783-A2
                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                             colloid derivatisation.
                                                                                                                                                                                                                                                                                                                                                                                               Alpha V beta 3 receptor; colloid-electrode binding; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alpha V beta 3 receptor-specific peptide
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(POLL/)
(ECHE/)
(MEAD/)
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17-SEP-1999;
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POLLOCK D.
ECHELARD Y.
MEADE H M.
RYBAK S M.
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ilarity 100.0%;
Conservative
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                                              99US-0116975P.
99US-0132289P.
99US-0133148P.
99US-0133772P.
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Pred. No. 0.0
0; Mismatches
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0.014;
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(MINE-) MINERVA BIOTECHNOLOGIES

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RESULT 15
AA022543
ID AA022
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Use of a composition comprising L-histidine, quisqualic acid, cycloserine, and suramin or their analogs for the treatment c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified.
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Patent No. 5534254
GENERAL INFORMATION:
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                                                                                                                                                                                         TELEFAX: 617-240 INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS: 1.ENGTH: 15 amino acids
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40, App	Sequence	US-08-599-654-40	N	1481		54	9
40,	Sequence	US-08-616-844-40	N	1481	81.8	54	8
•	Sequence	US-09-826-752-14	4	475		54	7
-	Sequence	US-09-323-433A-14	w	475		54	0
•	Sequence	US-08-396-001-14	N	475		54	5
14, Appl	Sequence	US-08-861-464-14	N	475		54	4
6442	Sequence	US-09-543-681A-6442	4.	408		54	ω
16550	Sequence	US-09-248-796A-16550	4	239		54	N
1, Appli	Sequence	US-08-257-341-1	ب	13	81.8	54	1
41907	Sequence	US-09-270-767-41907	4	212		57	0
16,	Sequence	-786-	N	622	90.9	60	φ
2, Appl	Sequence	US-08-461-386-2	N	250		60	8

ALIGNMENTS

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; LOCATION: 1..15; OTHER INFORMATION: /note= "LINKER US-08-133-804-8
                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/133
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08133804
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Place, 53 State
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Score 66;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thibeault/Patent Department
State Street
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DB 1;
0.0087;
              Length
                   15;
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RESULT 3
US-08-461-386-8
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                                                                         Sequence 8, Application US/08461386 Patent No. 5837846
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GENERAL INFORM
                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 15; Conserv
                                                             GENERAL INFORMATION:
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 APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 617-248-7100 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,838
FILLING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                    NAME/KEY: Protein
LOCATION: 1..15
COTHER INFORMATION: /note= "LINKER 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Kelley, Robin D. REGISTRATION NUMBER: 34 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: M
                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
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Exchange Place, 53 State Street
Oppermann, Hermann
Houston, L. L.
Ring, David B.
                                           Huston, James S
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Houston, L. L
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                                                                                                                                                                                                                                          100.0%; Score 66; DB 1; 100.0%; Pred. No. 0.0087;
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US-08-356-786-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Local Similarity
Conser
                                                                                                                                                                                                                                                                                                                                                    Sequence 12,
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                        NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                        APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Protein for Cancer
TITLE OF INVENTION: Marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                             COUNTRY:
                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: Protein LOCATION: 1..15
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OTHER INFORMATION: /note= "LINKER 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 424
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                                                                                                02109
                                                                                                                                             Boston
                                                                                                                                                                                                                                                                                                                                       2, Application US/08356786
5877305
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7: USA
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Houston, L.
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100.0%; ;
htive 0;
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Imaging
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                Version
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                #1.25
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 424

US/08/356,786

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MOLECULE TYPE: protein
US-08-356-786-12
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                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PAtentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/665,202

PILING DATE: 13-UN-1996

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/000,238

FILING DATE: 14-UN-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/000,250

FILING DATE: 14-UN-1995

PRIOR APPLICATION USBER: US 60/000,250

PRIOR APPLICATION NUMBER: US 60/000,250
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                                         TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Marks, Ja
APPLICANT: Schier, R
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Pitcher, Edmund R.
REGISTRATION NUMBER: 27,829
REFERENCE/DOCKET NUMBER: CRP-053
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
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FILING DATE: 06-FEB-1992
ATTORNEY/AGENT INFORMATION:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                          REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 02:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                     FILING DATE: 15-JUN-1995
ATTORNEY/AGENT INFORMATION:
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Local Similarity 100.0%; (
hes 15; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Antigens
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; Pred. No. 0.00
0; Mismatches
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                                                                                                                    US-09-315-574-3
                                                                                                                                  RESULT 7
                                                                                                                                                                                                                                                                                                                      US-08-483-749A-31
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                                                                                                                                                                                                                                                    Best Loc
Matches
                                                                                 Sequence 3, Application US/09315574 Patent No. 6512097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94662-8097
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPATIBLE: THEM PC COMPATIBLE
COMPATIBLE: TYPEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
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Best Local Similarity
                                                                                                                                                                                                                                                                                   Query Match
                                 GENERAL INFORMATION:
APPLICANT: Marks, James D.
APPLICANT: Schier, Robert
                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (510) 655-3542
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/483,749A
FILING DATE: 07-7UN-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: RING, DAVID B.
TITLE OF INVENTION: ANTIGEN-BINDING SITES OF ANTIBODY
TITLE OF INVENTION: MOLECULES SPECIFIC FOR CANCER ANTIGENS
APPLICANT: Schier, Robert TITLE OF INVENTION: No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                ILECULATION (510)
                                                                                                                                                                                                                                                                                                                                                   LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                  Local Similarity es 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: SAVEREIDE, PAUL B
REGISTRATION NUMBER: 36
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: CHIRON CORPORATION
STREET: INTELLECTUAL PROPERTY - R440, PO BOX 8097
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                                                                                                                                                                                                                                                                                                                                ESS: single
linear
                                                                                                                                                                                                                                                    100.0%; Score 66; DB 3; ilarity 100.0%; Pred. No. 0.0087; Conservative 0; Mismatches 0
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 Tumor Antigens
                 No. 6512097el High Affinity Human Antibodies to
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Pred. No. 0.0087;
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; OTHER INFORMATION: linker US-09-324-258-18
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US-09-324-258-18
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                                                                                                       SOFTWARE: F
SEQ ID NO 18
LENGTH: 15
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GENERAL INFORMATION:
                                                                                                                                                               APPLICANT: Bandad, Cynthia C.

TITLE OF INVENTION: THE USE OF SELF-ASSEMBLED MONOLAYERS TO TITLE OF INVENTION: PROBE THE STRUCTURE OF A TARGET MOLECULE FILE REFERENCE: M1015/7004/TJO CURRENT APPLICATION NUMBER: US/09/324,258
CURRENT FILING DATE: 1999-06-02
PRIOR APPLICATION NUMBER: US. 60/087,766
PRIOR FILING DATE: 1998-06-02
RIVERENT FILING DATE: 1998-06-02
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Best Local Similarity
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APPLICATION NUMBER: US 60/000,250
FILING DATE: 15-UUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/665,202
FILING DATE: 13-UUN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids
                                              FEATURE:
                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptide
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REFERENCE/DOCKET NUMBER: 02307E-061411
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION UMBER: US/09/315,574 FILING DATE: 20-MAY-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,238
FILING DATE: 14-UN-1995
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                                                                                                                                                   FastSEQ for Windows Version 3.0
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                ; OTHER INFORMATION: Linker sequence US-10-077-210-4
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US-09-333-213-4
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Best Local S
Matches 15
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APPLICANT: Weade, Harry
APPLICANT: KTANE, ISONOFITIN ANALOGY-HUMAN SERUM ALBUMIN FUSION
FILE REFERENCE: 10275/041001
CURRENT APPLICATION NUMBER: US/09/333,213
CURRENT FILING DATE: 1999-06-15
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 17
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                          Query Match
                                                                                                                                                                                                               APPLICANT: ML, COUNG-Hwa
APPLICANT: Lin, Rong-Hwa
APPLICANT: Hsu, Pei-Ling
TITLE OF INVENTION: PRODUCTION OF RECOMBINANT PROTEINS IN VIVO AND USE FOR GENERATING
TITLE OF INVENTION: ANTIBODIES
FILE REFERENCE: 13062-004001
CURRENT APPLICATION NUMBER: US/10/077,210
CURRENT FILING DATE: 2002-06-13
FRIOR APPLICATION NUMBER: 60/318,474
PRIOR PILING DATE: 2001-09-10
NUMBER OF SEO ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local
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                                                                                                                                                LENGTH: 17
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Synthetically generated linker sequence
                                                          Local
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                                                       Similarity
   SSSSGSSSSGSSSSG 15
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                                    100.0%; Score 66; DB 4; Length 17; 100.0%; Pred. No. 0.0099; Live 0; Mismatches 0; Indels
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Pred. No. 0.
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NESULT 13
US-08-1804-6
; Sequence 6, Application US/08133804
; Patent No. 5534254
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US-09-324-258-19
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/411,067C CURRENT FILING DATE: 2002-05-07 NUMBER OF SEQ ID NOS: 5 SOPTWARE: Patentin version 3.0 SEQ ID NO 5 LENGTH: 100 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09411067C
Patent No. 6576610
GENERAL INFORMATION:
APPLICANT: NUVAS, LLC
APPLICANT: HOUSTON, L.L.
APPLICANT: HOUSTON, L.L.
TITLE OF INVENTION: USE OF A CONTEXT-DEPENDENT FUNCTIONAL ENTITY TO ENHANCE THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 19
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Patent No. 6723517
GENERAL INFORMATION:
APPLICANT: Bamdad,
                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/324,258
CURRENT FILING DATE: 1999-06-02
PRIOR APPLICATION NUMBER: U.S. 60/087,766
PRIOR FILING DATE: 1998-06-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: NUVAS1140
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                                                                                                                                                                                                                                                                                                              NAME/KEY: REPEAT
LOCATION: (1)..(5)
OTHER INFORMATION: Spacer peptide variation; sequence repeated "n" times, where n =
OTHER INFORMATION: 1-20
                                                                                                                                                                                                                                                                                                                                                                                                 PEATURE:
OTHER INFORMATION: Spacer element
PEATURE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANIM: Artificial Sequence
PEATURE:
OTHER INFORMATION: 1X-linker-1X
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15; Conser
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                                                                                                                                                                                                                             DB 4; Length 100; 0.062;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION: APPLICANT: Huston
                                                                                                                                                                                                                                 APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For
TITLE OF INVENTION: Imaging
             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 243 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Kelley, Robin D. REGISTRATION NUMBER: 34, REFERENCE/DOCKET NUMBER:
                                                                                                                COUNTRY: U
ZIP: 02109
                                                                                                                                                                     CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 424
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STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 617-248-7477
TELEFAX: 617-248-7100
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State Street
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##NGTH: 243 amino acids
TYPE: amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-461-838-6
                                                                                    INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS: LENGTH: 243 amino acids TYPE: amino acids TYPE: amino acids TYPE: Information acids TYPE: TYPE: protein US-08-461-386-6
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US-08-461-386-6
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Query Match 100.0%; Score 66; DB 2; Length 243; Best Local Similarity 100.0%; Pred. No. 0.16; Matches 15; Conservative 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,386
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REFERENCE/DOCKET NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 6, Application US/08461386 Patent No. 5837846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 66; DB 1
Best Local Similarity 100.0%; Pred. No. 0.16;
Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Herma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 243 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREE: Boston
CITY: Boston
STATE: Massachusetts
THINTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
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FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 32,637
REFERENCE/DOCKET NUMBER: 2054/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Testa, Hu
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Qy 1 SSSGSSSSGSSSSG 15
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Db 118 SSSGSSSSGSSSSG 132
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Search completed: April 4, 2005, 16:00:54 Job time: 8.66773 secs

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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2=6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2-6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

3: /cgn2-6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

4: /cgn2-6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

6: /cgn2-6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

6: /cgn2-6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

7: /cgn2-6/ptodata/1/pubpaa/US08 NEW PUB.pep:*

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17: /cgn2-6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

18: /cgn2-6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

19: /cgn2-6/ptodata/1/pubpaa/US10 PUBCOMB.pep:*

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US-10-204-581-1
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7 US-10-683-547-8
3 US-10-081-400-4
1 US-10-077-213-4
1 US-10-077-210-4
1 US-10-078-873-4
US-10-768-873-4
US-10-996-069-12
US-10-003-681-1
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RESULT 1

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US-09-840-746-1	10-282-122	-10-108-	10-108-605-12	10-258-951	10-032-585	10-912-434-1	US-10-369-493-22092	9-826-752-1	-10-25	-10-108-2607	10-451-4671	98-512-	-10-050-704-	-10-156-761-119	0-264	-10-437-963-19922	-10-424-599	US-09-925-301-1154	09-764-864-	-09-764-864-	9-764-864-	-09-764-864-12	10-437-963-1985	10-767-701-3	10-369-493-609	10-683-547-	φ	-10-683	9-887-853-	-10-683-54	US-09-887-853-6
Sequence 1, Appli	equence 782	e 129,	e 125,	e 72,	e 7924	ė	e 2209	4	73, 1	ە ھ	262,	e 100,	e 100,	e 11	e 1938,	e 1992	ce 1756	1154,	812,	817,	_	1273,	ø	e 35941	ē O	е 2	Ŋ	e 4	4	Sequence 6, Appli	Sequence 6, Appli

ALIGNMENTS

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RESULT 2
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; Sequence 8, Application US/09887853
; Patent No. US20020168375A1
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                                                                                                                                                                                                                              Query Match
Best Local S
Matches 15
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APPLICANT: Amersdorfer, Peter
APPLICANT: Amersdorfer, Peter
TITLE OF INVENTION: Therapeutic Monoclonal Antibodies That Neutralize
TITLE OF INVENTION: Botulinum Neurotoxins
FILE REFERENCE: 2500.117USO
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT APPLICATION NUMBER: US/09/144,886
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3
LENGTH: 15
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
TEATURE:
OTHER INFORMATION: Description of Artificial Sequence: [(Ser)4Gly]3
OTHER INFORMATION: linker
                                                                                                                                                                                                                                l Similarity
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RESULT 3
US-10-204-581-1
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FILING DATE: <UNKNOWN:
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7100
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Zardi, Luciar
APPLICANT: Neri, Dario
                                                                                                                                                                                                                                  Sequence 1, Application US/10204581 Publication No. US20040013640A1
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
            APPLICANT: Borsi, Laura
APPLICANT: Halin, Cornelia
TITLE OF INVENTION: On pathological lesions
TITLE OF INVENTION: in pathological lesions
FILE REFERENCE: 0380-P02964US0
                                                                                                                      APPLICANT:
                                                                                                                                                       APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/09/887,853
FILING DATE: 21-Jun-2001
CLASSIFICATION: <UNKnown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 1..15
OTHER INFORMATION: /note= "LINKER 2"
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US/10/204,581
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                  Carnemolla, Barbara
Nilsson, Fredrik
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                                                                                                              Tarli, Lorenzo
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INVENTION: Biosynthetic Binding Proteins For
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PRIOR APPLICATION NUMBER: PCT/IB01/00382
PRIOR FILLING DATE: 2001-02-22
PRIOR FILLING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: US 60/184,767
PRIOR APPLICATION NUMBER: US 60/257,192
PRIOR FILLING DATE: 2000-12-21
PRIOR FILLING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
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; OTHER INFORMATION: Description of Artificial Sequence: Linker US-10-204-581-1
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Best Local S
Matches 15
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TITLE OF INVENTION: THERAPBUTIC MONOCLONAL ANTIBODIES THAT NEUTRALIZE BOTULINUM
TITLE OF INVENTION: NEUROTOXINS
FILE REFERENCE: 407T-895120US
FILE REFERENCE: 407T-895120US
CURRENT APPLICATION NUMBER: US/10/632,706
CURRENT FILING DATE: 2003-08-01
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 60/400,721
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR APPLICATION NUMBER: US 09/144,806
PRIOR FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 278
SOFTWARE: Patentin version 3.2
SEQ ID NO 3
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Publication No. US20040175385A1
GENERAL INFORMATION:
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Publication No. US20050058638A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 15; Conserv
APPLICANT: HOUSTON, L.L.
APPLICANT: Ring, D.
APPLICANT: Oppermann, H.
TITLE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING
FILE REFERENCE: CIBT-P01-130
CURRENT APPLICATION NUMBER: US/10/683,547
                                                                                                                             APPLICANT: Huston, J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: protein linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial FEATURE:
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100.0%; Pred. No. 0.062;
tive 0; Mismatches
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CURRENT FILING DATE:

2003-10-10

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GENERAL INFORMATION:

APPLICANT: Heddey, MAIY Lynne
ITITLE OF INVENTION: METHODS OF TREATING BLADDER DISORDERS
ITILE REFERENCE: 08191-022001

CURRENT APPLICATION NUMBER: US/10/074,956

CURRENT FILING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: 600/268,175
PRIOR FILING DATE: 2001-02-12

NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 19
LENGTH: 17
                                                                                                                                                                                                                                                   US-10-074-956-19
; Sequence 19, Application US/10074956
; Publication No. US20020193332A1
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US-10-081-400-4
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LENGTH: 17
TYPE: PRT
ORGANISM: Artificial Sequence
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SEQ ID NO 8
LENGTH: 15
TYPE: PRT
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Best Local Similarity 100.0%;
Matches 15; Conservative (
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APPLICANT: Meade, Harry
APPLICANT: Meade, Harry
APPLICANT: Krane, Ian
TITLE OF INVENTION: ERYTHROPOIETIN ANALOGY-HUMAN SERUM ALBUMIN FUSION
FILE REFERENCE: 10275/041001
CURRENT APPLICATION NUMBER: US/10/081,400
CURRENT FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 09/333,213
PRIOR PILING DATE: 1999-06-15
PRIOR FILING DATE: 1999-06-15
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PRIOR FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 07/831,967
PRIOR FILING DATE: 1992-02-06
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 4.0
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tes 15; Conservation
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Pred. No. 0.062;
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Pred. No. 0.069;
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                                 ; OTHER INFORMATION: Linker sequence US-10-077-210-4
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                                                                                                                                                                                                                                                                  Sequence 4, Application US/10077210
Publication No. US20030049801A1
GENERAL INFORMATION:
APPLICANT: Wu, Chung-Hsiun
APPLICANT: Lin, Rong-Hwa
APPLICANT: Lin, Rong-Hwa
APPLICANT: Hsu, Pei-Ling
TITLE OF INVENTION: PRODUCTION OF RECOMBINANT PROTEINS
TITLE OF INVENTION: ANTIBODIES
FILE REFERENCE: 13062-004001
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Best Local Similarity
Matches 15; Conserva
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Publication No. US20030049694A1
GENERAL INFORMATION:
                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4
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Query Match
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                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/077,210 CURRENT FILING DATE: 2002-06-13 PRIOR APPLICATION NUMBER: 60/318,474 PRIOR FILING DATE: 2001-09-10 NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lin, Rong-Hwa
APPLICANT: Hau, Pei-Ling
TITLE OF INVENTION: PRODUCTION OF FUSION PROTEINS AND USE
TITLE OF INVENTION: POR IDENTIFYING BINDING MOLECULES
FILE REFERENCE: 13062-002001
CURRENT APPLICATION NUMBER: US/10/077,213
CURRENT FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: 60,318,474
PRIOR PILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 17
TYPE: PRT
ORGANISM: Artificial :
FEATURE:
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APPLICANT: Lin, Rong-Hwa
APPLICANT: Hsu, Pei-Ling
                                                                                    LENGTH: 17
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                       FEATURE:
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100.0%; Score 66;
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Pred. No. 0.069;
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Pred. No. 0.069;
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Length 17;
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Best Local Similarity

100.0%; Pred. No. 0.069;

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RESULT 11
US-10-768-873-4
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; ORGANISM: peptide linker
US-10-608-710-9
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US-10-608-710-9
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                                                                                                                                     US-10-768-873-4
                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Young, Michael
APPLICANT: Weade, Harry
APPLICANT: Krane, Ian
TITLE OF INVENTION: ERYTHROPOIETIN ANALOGY-HUMAN SERUM ALBUMIN FUSION
FILE REFERENCE: 10275/041001
CURRENT APPLICATION NUMBER: US/10/768,873
CURRENT FILING DATE: 1094-01-30
PRIOR APPLICATION NUMBER: US/09/333,213
PRIOR APPLICATION NUMBER: US/09/333,213
PRIOR FILING DATE: 1999-06-15
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:

APPLICANT: GTC Biotherapeutics, Inc.
APPLICANT: Edge, Michael D
APPLICANT: Pollock, Daniel
APPLICANT: Echelard, Yann
APPLICANT: Meade, Harry M
APPLICANT: Rybak, Susanna M
TITLE OF INVENTION: Transgenically Produced Fusion Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.2
SEQ ID NO 9
LENGTH: 17
                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4
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                                                                  Matches
                                                                                   Best Local Similarity
                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: GTC-42D
CURRENT APPLICATION NUMBER: US/10/608,710
CURRENT FILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: US 09/398,610
PRIOR FILING DATE: 1999-09-17
                                                                                                                                                                                    LENGTH: 17
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                    OTHER INFORMATION: Synthetically generated linker sequence
                                                                                                                                                                       FEATURE:
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                                                                 Conservative
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                                                                                 100.0%; Score 66; DB 16; 100.0%; Pred. No. 0.069;
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Pred. No. 0.069;
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; Sequence 6, Application US/09887853 ; Patent No. US20020168375A1 ; GENERAL INFORMATION:
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                                                                             RESULT 14
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                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Synthetic sequence US-10-003-681-1
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US-10-003-681-1
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                                                          US-09-887-853-6
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APPLICANT: Bamdad, R. Shoshana
APPLICANT: Bamdad, R. Shoshana
TITLE OF INVENTION: DIAGNOSTIC TUMOR MARKERS, DRUG SCREENING FOR TUMORIGENESIS INHIBI7
TITLE OF INVENTION: AND COMPOSITIONS AND METHODS FOR TREATMENT OF CANCER
TILE REPERENCE: M01015/70071
CURRENT APPLICATION NUMBER: US/09/996,069
CURRENT FILING DATE: 2001-11-27
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin version 3.1
SEQ ID NO 12
LENGTH: 31
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/10003681
Publication No. US20020156112A1
GENERAL INFORMATION:
APPLICANT: Bandad, R. Shoshana
APPLICANT: Bandad, Cynthia
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Publication No. US20030036199A1
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Best Local Similarity
Matches 15; Conserva
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/248,865
PRIOR FILING DATE: 2000-11-15
PRIOR APPLICATION NUMBER: US 60/277,922
PRIOR FILING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Bamdad, Cynthia
TITLE OF INVENTION: ENDOSTATIN-LIKE ANGIOGENISIS INHIBITION
FILE REFERENCE: M01015.70046.US
CURRENT APPLICATION NUMBER: US/10/003,681
CURRENT FILING DATE: 2001-11-15
                                                                                                                                                                                                                                                                                                                                                     LENGTH: 31
TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                     FEATURE:
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100.0%; Pred. No. 0.12;
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Pred. No. 0.12;
Nismatches
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APPLICANT: Huston, James

Oppermann, Hermann Houston, L. L.

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Sequence 6, Application US/10683547

Publication No. US20050058638A1

GENERAL INFORMATION:
APPLICANT: Houston, J.
APPLICANT: Ring, D.
APPLICANT: Ring, D.
APPLICANT: Oppermann, H.
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US-10-683-547-6
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                                                                                                              APPLICANT: ODDERMAIN, H.

TITLE OF INVENTION: BIOSYNTHETIC BINDING PROTEINS FOR IMMUNO-TARGETING FILE REFERENCE: CIBT-P01-130
CURRENT APPLICATION NUMBER: US/10/683,547
CURRENT FILING DATE: 2003-10-10
PRIOR APPLICATION NUMBER: US/99/558,741
PRIOR FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 07/831,967
PRIOR APPLICATION NUMBER: 07/831,967
PRIOR FILING DATE: 1992-02-06
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.1
150 ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/13,804
FILING DATE: <UNKNOWN'
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 2054/22
REFERENCE/DOCKET NUMBER: 2054/22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7100
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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Matches 15; Conserv
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                                         LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
OTHER INFORMATION: 520C9 sev
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/887,853

FILING DATE: 21-Jun-2001

CLASSIFICATION: CUnknown>

PRIOR APPLICATION DATA:

CLASSIFICATION DATA:

PRIOR APPLICATION DATA:
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
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ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 11
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TITLE OF INVENTION: Biosynthetic Binding Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       118 sssssssssssss 132
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STATE: Massachusetts
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Search completed: April 4, 2005, 16:42:47 Job time : 23.1406 secs
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Best Local Similarity
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                                                                            1 SSSSGSSSSGSSSSG 15
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       nuclear
                                                                                                                                                                                                                                 hypothetical
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	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30
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	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8	75.8
	1732	1723	1460	1446	1029	772	742	741	712	710	625	559	440	419	380
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ALIGNMENTS	C81601	H86557	EDBEIF	A45344	T30351	A55004	A49672	I48694	B47021	S28014	T41603	RWBYS1	S37303	T14448	T14447
	polymorphic membra		immediate-early pr	immediate-early pr	mucin-like protein	transcription fact	transcription fact	probable transcrip	pectic enzyme secr	outD protein - Erw	alpha-amylase - fi	glycophospholipid-	sox-4 protein - mo	oleosin homolog pc	oleosin homolog pc

R/Bentley, D.

R/Bent

The sequence

cosmid

K02F3

A; Reference number: A; Accession: T16535

A;Status: preliminary; translated from GB/EMBL/DDBJ

hypothetical protein K02F3.11 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 20-Sep-1999 C;Accession: T16535

T16535

RESULT 1

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RESULT 2
T26294
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A;Molecule type: DNA
A;Residues: 1-304 <BEN>
A;Cross-references: EMBL:U00052; NID:g485125; PID:g485126; PIDN:AAA50702.1; CESP:K02F3.1]
A;Cross-references: EMBL:U00052; NID:g485125; PID:g485126; PIDN:AAA50702.1; CESP:K02F3.1]
A;Map position: 4
A;Map position: 4
A;Introns: 32/3; 58/3; 117/3; 211/3; 263/3; 675/3; 1139/3; 1187/1; 1230/3
A;Introns: 32/3; 58/3; 117/3; 211/3; 263/3; 675/3; 1139/3; 1187/1; 1230/3
C;Superfamily: Na+/K+-transporting Arrase alpha chain; ATPase nucleotide-binding formaln homology <ATN>
                                                                                                                                                                                              A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                    A; Reference number: Z20191
A; Accession: T26294
                                                                                                                                                                                                                                                                                                                hypothetical protein W09C2.3 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 18-Aug-2000
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A; Introns: 257/2
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                                                                                                                                                                                                                                                           submitted to the EMBL Data Library, December 1995
                                                                                                                                                                                                                                                                                                C; Accession: T26294
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Pred. No. 0.98;
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Query Match

86.4%;

Score 57;

DB

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Length 1249;

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A;Experimental source: strain 9a5c

A;Experimental source: strain 9a5c

A;Simpson, A.J.G.; Reinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; A Briones, M.R.S.; Bueno, M.R.P.; Camargo, A.A.; Camargo, L.B.A.; Carraro, D.M.; Carrer, H. Briones, M.R.S.; Bueno, M.R.P.; Feaningon, A.P.; Ferreira, A.J.S.

as-Neto, E.; Docena, C.; El-Dorry, H.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohm Submitted to GenBank, June 2000

A;Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohm J.D.; Junqueira, M.L.; Kemper, B.L.; Kltajima, J.P.; Krieger, J.E.; Kuramae, E.E.; Laigrando, M.A.; Madeira, A.M.B.N.; Madeira, H.M.F.; Marino, C.I.; Marques, M.V.; Martins, E.A; Authors: Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, B.C.; Miyaki, C.Y.; F.G.; Numes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.A.; A.J. de M.; de Rosa Jr., V.B.; de Sa, R.G.; Santelli, R.V.; Sawasak A; Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva Jr., W.A.; da Silveira, M.; Tsuhhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore, A.L.; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1,4-beta-cellobiosidase XF1267 [imported] - Xylella fastidiosa (strain 9a5c: C;Species: Xylella fastidiosa
C;Species: Xylella fastidiosa
C;Date: 18e-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
C;Accession: A82704
R;anonymous, The Xylella fastidiosa Consortium of the Organization for Nucle
Nature 406, 151-157, 2000
A;Title: The genome sequence of the plant pathogen Xylella fastidiosa.
A;Reference number: A82515; MUID:20365717; PMID:10910347
A;Rote: for a complete list of authors see reference number A59328 below
A;Accession: A82704
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPROT:Q9PDW2; GB:AE003960; GB:AE003849; NID:g9106242; PIDN:AAF840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-683 <S
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A; Residues: 1-1751 <GHS-
A; Cross-references: UNIPROT: Q26194; GB: M75674; NID: g160608;
A; Note: sequence extracted from NCBI backbone (NCBIN: 83591, C; Superfamily: major merozoite surface antigen
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A;Accession: A45604
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C;Species: Plasmodium vivax
C;Date: 22-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
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                                 Query Match
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  Score 56; DB
Pred. No. 6.4;
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1; Mismatches
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NCBIP:83592)
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C;Accession: S38170; S40645; S3770
R;Baladron, V.; Ballesta, J.P.G.;
submitted to the Protein Sequence
                                                                                 SRP40 protein - yeast (Saccharomyces cerevisiae) N;Alternate names: protein YKR092c; protein YKR412a C;Species: Saccharomyces cerevisiae
                                                                                                                                                  RESULT
S38170
                                        C;Date: 03-May-1994 #sequence revision 03-May-1994 #text_change 09-Jul-2004 C;Accession: S38170; S40645; S37702
                                                                                                                                                                                                                                                                                                                                                                                      A;Gene: AT4g30460
A;Map position: 4
C;Superfamily: glycine-rich cell
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R; anonymous, The Eur
Nature 402, 769-777,
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A; Residues: 1-162 <STO>
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 16-Peb-2001 #sequence_revision 16-Peb-2001 #text_change 09-Jul-2004
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A; Map position: 4
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A;Title: Sequence and analysis of chromosome 4 of the plant Arabidopsis thaliana.
A;Reference number: A85001; MUID:20083488; PMID:10617198
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004
C;Accession: B85356
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86.7%;
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      Bou, G.;
Database,
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      G.; del Rey, F.
base, March 1994
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                       F.; Esteban,
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                     P.F.; Garcia-Cantalejo,
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probable membrane protein YIL123w - yeast (Saccharomyces cerevisiae)
N;Alternate names: hypothetical protein YI8277.06
C;Species: Saccharomyces cerevisiae
C;Date: 13-Jan-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
C;Accession: 849886
R;Hamlyn, N.; Churcher C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Robert, L.S.; Gerster, J.; Allard, S.; Cass, L.; Simmonds, J. Plant J. 6, 927-933, 1994
A;Title: Molecular characterization of two Brassica napus genes A;Reference number: 216348; MUID:95152562; PMID:7849761
A;Accession: T08093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: DNA
A;Residues: 1-406 <BOU>
A;Residues: 1-406 <BOU>
A;Cross-references: EMBL:X73541; NID:g450550; PIDN:CAA51946.1; PID:g450552
A;Experimental source: strain S288C
R;Lalo, D.; Carles, C.; Sentenac, A.; Thuriaux, P.
submitted to the EMBL Data Library, May 1993
A;Description: Interactions between three common subunits of yeast RNA polymerases I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-424 <ROB>
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A;Residues: 1-399,'N',401-406 <LAL>
Ā;Cross-references: EMBL:L11275; NI
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R;71tle: The complete sequence of a 15 820 bp segment of Saccharomyces cerevisiae chromc A;Reference number: $40644; MUID:94205265; PMID:8154186
A;Accession: $40645
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A; Molecule type: DNA
A; Residues: 1-406 <Bi
A; Cross-references: 1
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A;Accession: 837702
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;Species: Brassica napus (rape)
;Date: 21-May-1999 #sequence_revision 21-May-1999 #text_change 09-Jul-2004
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;Experimental source: cv. Westar; tapetum
;Note: protein component of lipid storage bodies of plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Accession: T08093
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Experimental source: strain S288C
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Best Local !
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Pred. No. 6.9;
0; Mismatches
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A;Cross-references: UNIPROT:P51022; EMBL:X69166; NID:g288079; PIDN:CAA48916.1; PID:g28808 R;Chan, T.; Bunting, M.; Karim, F.D.; Thummel, C.S.
Dev. Biol. 151, 176-191, 1992
A;Title: Isolation and Characterization of five Drosophila genes that encode an ets-related A;Reference number: $28819; MUID:92249640; PMID:1577186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene pointed protein splice form 1 - fruit fly (Drosophila melanogaster)
N;Alternate names: ETS-like protein
C;Species: Drosophila melanogaster
C;Date: 13-Jan-1995 #sequence revision 30-May-1997 #text_change 09-Jul-2004
C;Accession: S33167; S28819; S28818
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A;Reference number: S49881
A;Accession: S49886
A;Molecule type: DNA
A;Residues: 1-475 <HAM>
A;Residues: 1-475 <HAM>
A;Cross-references: UNIPROT:P40472; GB:Z47047; EMBL:Z46833; NID:g603997; PID:g763223; MII
                                                                                            C;Superfamily: pointed protein, splice form 1; etc C;Keywords: alternative splicing; DNA binding; nuc F;517-595/Domain: ets DNA-binding domain homology
                                                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 445-466, 'R', 468-603 <PRI>
A; Residues: 445-466, 'R', 468-603 <PRI>
A; Cross-references: EMBL.M20408; NID:g157384; PIDN:AAA28521.1; PID:g157385
A; Note: the authors translated the codon AGA for residue 467 as Gln
                                                                                                                                                                                                                                                                                                                                                            A;Title: The Drosophila ets-2 gene: molecular structure, A;Reference number: $28818; MUID:88196618; PMID:2834248 A;Accession: $28818 a;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:M88472; NID:g157191; PIDN:AAC34200.1; PID:g157192 R;Pribyl, L.J.; Watson, D.K.; McWilliams, M.J.; Ascione, R.; Papas, T.S. Dev. Biol. 127, 45-53, 1988
                                                                                                                                                                  A;Cross-references: FlyBase:FBgn0003118
A;Introns: 550/3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 456-613 < CHE>
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A;Title: The Drosophila gene pointed encodes two ETS-like
A;Reference number: 833167; MUID:94038653; PMID:8223245
A;Accession: 833167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Keywords: transmembrane protein
F;82-98/Domain: transmembrane #status predicted
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C;Species: Homo sapiens (man)
C;Decies: Homo sapiens (man)
C;Dete: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
C;Decession: 152882
C;Accession: 152882
R;Bloch, D.B.; Rabkina, D.; Quertermous, T.; Bloch, K.D.
Clin, Immunol: Immunopathol. 72, 380-389, 1994
A;Title: The immunoreactive region in a novel autoantigen contains a nuclea
A;Reference number: 152882; MUID:94340813; PMID:7520377
A;Cross-references: UNIPROT:060382; C;Genetics:
                                           A; Molecule type: DNA
A; Residues: 1-1791 <RIC>
                                                                                                                          A; Description: Sequencing of A; Reference number: Z14664
                                                                                                                                                         C;Accession: T02345

R;Ricke, D.O.; Bruce, D.; Mundt, M.; Doggett, N.; Munk, C.; Saunders, E.; Robinson, re, J.; White, S.; Ueng, S.; Tatum, O.; Campbell, C.; Fawcett, J.; Deaven, L. submitted to the EMBL Data Library, March 1998
                                                                                                                                                                                                                                          hypothetical protein KIAA0324 - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 05-Mar-1999 #sequence_revision 05-Mar-1999 #text_change 09-Jul-2004
                                                                                 A; Status: preliminary;
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A;Cross-references: UNIPROT:Q13826; GB:L26339; NID:g533201; PIDN:AAA21833.1; PID:g533202
C;Superfamily: human autoantigen Ge-1
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A;Status: preliminary; translated from GB/EWBL/DDBJ
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A; Residues: 1-679 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barro, Nature 413, 523-527, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probable membrane protein YPO2731 [imported] - Yersinia pestis (strain CO92)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;Species: Yersinia pestis;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001;Accession: AC0333
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0; Mismatches
                 EMBL: AC004493; NID: g2996648; PIDN: AAC08453.1;
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S.; Barrell,
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neutrophil protein - pig (fragment)
(;Species: Sus scrofa domestica (domestic pig)
(;Species: 18-Jan-1991 #sequence_revision 18-Jan-1991 #text_change 09-Jul-2004
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A; Residues: 1628-1639 < YAM>
A; Residues: 1628-1639 < YAM>
C; Superfamily: vitellogenin
C; Keywords: egg yolk; glycoprotein; liver; phosphoprotein
C; Keywords: egg yolk; glycoprotein; phosphoprotein
C; Keywords: egg yolk; glycoprotein; liver; phosphoprotein
C; Keywords: egg yolk; glycoprotein; liver; phosphoprotein
C; Keywords: egg yolk; glycoprotein; liver; phosphoprotein
                                                                                                                                                                                                                 Biochem. Biophys. Res. Commun. 170, 915-922, 1990
A;Title: Isolation and characterization of a cDNA clone
A;Reference number: A35419; MUID:90343818; PMID:1696479
                                                                                                                                                                                                                                                                                  C;Accession: A35419
R;Bellavite, P.; Bazzoni, F.; Cassatella, M.A.; Hunter, K.J.; Bannister, J.V.
                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-284 <BEL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Experimental source: liver
R;Yamamura, J.; Adachi, T.; Aoki, N.; Nakajima, H.; Nakamura, R.; Matsuda,
Biochim. Biophys. Acta 1244, 384-394, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vitellogenin I precursor [validated] - chicken
N;Contains: lipovitellin I; lipovitellin II; phosvitin; yolk glycoprotein 42K
C;Species: Gallus gallus (chicken)
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T29088; S55681
R;Mabuchi, N.; Yamamura, J.; Adachi, T.; Aoki, N.; Nakamura, R.; Matsuda, T.
submitted to the EMBL Data Library, November 1996
                                                                                          A; Cross-references: UNIPROT: Q99331; GB: M55701; NID: g164672;
                                                                                                                                                                      A;Status: preliminary
                                                                                                                                                                                          A; Accession: A35419
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A;Accession: T29088
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A; Introns: 1610/2;
A; Note: KIAA0324
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86.7%; Pred. No. 8.6,
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Q6c6n1 yarrowia li
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Q9w4n4 drosophila
Q759n0 ashbya goss
Q7p947 anopheles g
Q8tq10 methanosarc
Q693b7 microbulbif
Q7qda9 anopheles g
Q8tq10 dictyosteli
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Barnay S., Blanchin S., Beckerich J.M., Beyne E., Bleykasten C.,
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Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
Hantraye F., Hemnequin C., Jauniaux N., Joyet P., Kachouri R.,
Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
Nicaud J.M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
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Similar to tr[Q08294 Saccharomyces cerevisiae YOL155c.
ORFNames=YALIOE08008g;
Yarrowia lipolytica CLIB99.
                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM STRAIN=CLIB99;
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Eukaryota; Fungi; Ascomycota;
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PROSITE; PS00030; RRM_RNP_1; UNKNOWN_1.
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NCBI_TaxID=284591;
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CG15570-PA.
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                       ORFNames=CG15570;
                                                                                                                                                                                                                                                  InterPro; IPR009071; HMG-box.
InterPro; IPR009910; HMG 12 b
Pfam; PF00505; HMG box; T.
SMART; SM00398; HMG; 1.
PROSITE; PS50118; HMG_BOX_2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).
Eukaryota; Metazoa; Chordata; Caraniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetradontiformes;
Tetradontoidea; Tetraodontidae; Takifugu.
                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transcription Name=Sox11;
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genoscope;
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; CR382131; CAG79270.1; -.
SEQUENCE 453 AA; 43707 MW; 097090FA7AE15F50 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q6WNT4;
                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=31033;
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                                            (TrEMBLrel. 13, (TrEMBLrel. 13, (TrEMBLrel. 25,
                                                                                                                                                                                                                                        368 AA;
                                                                                          PRELIMINARY;
                                                                                                                                                                                            Conservative
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40719 MW;
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Pred. No. 6.8;
0; Mismatches
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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Golder J.D.,
RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Golder J.D.,
RA Button G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Brandon R.C., Rogers Y.H., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Mark H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Gabor G.L.,
RA Abril J.F., Agbayani A., An H.J., Andrews-Pfannkoch C., Baldwin D.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Shandari D., Bothakov S.,
RA Glerry J.M., Cawley S., Dahlke C., Davanport L.B., Davies F.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies F.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies F.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies F.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies F.,
RA Cherry J.M., Cawley S., Domes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
Davis R.J., Congrellista C.C., Ferraz C., Ferriexa S., Fleischmann W.,
Rosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Howland T.J., Wei M.H., Ibegwam C.,
RA Liu X., Nattei B., McIncoh T.C., McLeed M.P., McCherson D.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D. Lei Z.,
Liang Y., Lin X.,
RA Melson D.R., Nalson K.A., Nixon K., Nusskern D.R., McCherson D.,
RA Winske B.E., Sodira C.D., For S., Pollard J., Puri V., Resee M.G.,
RA Melson D.R., Nalson K., Namders R.D., Scheeler F., Shen H.,
RA Nalson K.A., Mixon K., Nusskern D.R., Shich H.,
RA Nalson S.M., Woodastry C., Morris J., Moshredi A.,
RA Nalson K.A., Sanders R.D., Scheeler F., Shen H.,
RA Shue B.C., Stanker S., Pans S., Palo Q., Zhao Q., Ne J.,
RA Yelne F., Zaveri J.S., Zhan M., Zhang S., Zhu X., Smith H.O.,
RA Sheng X.H., Woodast
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Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
Misra S., Crosby M.A., Mungall C.J.,
Hradecky P., Huang Y., Kaminker J.S.,
Smith C.D., Tupy J.L., Whitified E.J.
Bettencourt B.R., Celniker S.E., de
Harris N.L., Richter J., Russo S., S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE-22426070; PubMed-12537573;
Kaminker J.S., Bergman C.M., Kronmiller
Patel S., Frise E., Wheeler D.A., Lewis
                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                          Ashburner M., Celniker S.E.;
"The transposable elements of the a genomics perspective.";
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Ephydroidea;
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sa; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                     PubMed=12537572;
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                                       Mungall C.J., Matthews B.B., Campbell K.S.,
Kaminker J.S., Milburn G.H., Prochnik S.E.,
Whitfied E.J., Bayraktaroglu L., Berman B.P
iker S.E., de Grey A.D., Drysdale R.A.,
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Q759NO;
O5-JUL-2004 (TrEMBLrel. 2
O5-JUL-2004 (TrEMBLrel. 2
O5-JUL-2004 (TrEMBLrel. 2
                                                                                                                                               Q7PQ47; PRELIMINARY;
Q7PQ47; C1-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                           ENSANGP00000021699 (Fragment)
Name=ENSANGG00000019210,
Anopheles gambiae str. PEST.
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                                                      Eukaryota; Metazoa; Arthropoda;
Neoptera; Endopterygota; Dipter
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SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ascomycota;
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                                                          Diptera;
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Pred. No.
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Saccharomycotina; Saccetaceae; Eremothecium
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                                                          Hexapoda; Ins
a; Nematocera;
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RX MEDILINE=21929760; PubMed=11932238; DOI=10.1101/gr.223902;

RA Galagan J.E., Nusbaum C., Rey A., Endrizzi M.G., Macdonald P.,

RA Galegan J.E., Nusbaum C., Rey A., Endrizzi M.G., Macdonald P.,

RA FitzHugh W., Calvo S., Engels R., Smirnov S., Atnoor D., Brown A.,

RA Allen N., Naylor J., Stange-Thomann N., DeArellano K., Johnson R.,

Linton L., McEwan P., McKernan K., Talamas J., Tirrell A., Ye W.,

RA Linton L., McEwan P., Cann I., Graham D.E., Grahame D.A., Guss A.M.,

RA Zimmer A., Barber R.D., Cann I., Graham D.E., Grahame D.A., Guss A.M.,

RA Hedderich R., Ingram-Smith C., Kuettner H.C., Krzycki J.A.,

RA Hedderich R., Jingram-Smith C., Kuettner H.C., Krzycki J.A.,

RA Hedderich R., Jingram-Smith C., White R.H., de Macario E.C.,

RA Springer T.A., Umayam L.A., White O., White R.H., de Macario E.C.,

RA Springer T.A., Umayam L.A., White O., White R.H., Daulsen I.T.,

RA Ferry J.G., Jarrell K.F., Jing H., Macario A.J.L., Paulsen I.T.,

RA Pritchett M., Sowers K.R., Swanson R.V., Zinder S.H., Lander E.,

RA Metcalf W.W., Birren B.;

RT and physiological diversity.";

Genome Res. 12:532-542(2002).
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Q693B7;
Q693B7;
25-OCT-2004 (TrEMBLrel. 2
25-OCT-2004 (TrEMBLrel. 2
25-OCT-2004 (TrEMBLrel. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q8TQ10;
Q8TQ10;
01-JUN-2002
01-JUN-2002
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Submitted (APR-2003) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-2002 (TrEMBLrel. 21, 01-JUN-2002 (TrEMBLrel. 21, 01-OCT-2002 (TrEMBLrel. 22, Hypothetical protein MA1742. OrderedLocusNames=MA1742;
     Endo-chitinase.
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Archaea; Euryarchaeota; Methanomicrobia; Methanosarcinales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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InterPro; IPR008962; PapD-like.
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GO; GO:0005576; C:extracellular; IEA.
GO; GO:0016787; F:hydrolase activity; IE
GO; GO:0004553; F:hydrolase activity; hy
GO; GO:0004553; F:hydrolase activity, hy
GO; GO:0005975; P:carbohydrate metabolis
GO; GO:0005152; P:metabolism; IEA.
InterPro; IPR005123; Glyco_hydro_18.
InterPro; IPR001223; Glyco_hydro_18.
InterPro; IPR001223; Glyco_hydro_18AS.
Pfam; PF00704; Glyco_hydro_18; 1.
SMART; SM00495; ChtED3; 2.

Pfam; PF00704; Glyco_hydro_18; 1.
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Q7QDA9;
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE:
STRAIN-PEST;
Anopheles Genome Sequencing Consortium;
Anopheles Genome Sequence to the EMBL/GenBank/DDBJ databases
Submitted (MAR-2002) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
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EMBL; GO:0005576; C:extracellular; IEA.
GO: GO:0005576; F:carbohydrate binding; IEA.
GO: GO:0016787; F:hydrolase activity, IEA.
GO: GO:0016787; F:hydrolase activity, Mydro
GO: GO:0004553; F:hydrolase activity, Mydro
GO: GO:0005975; P:carbohydrate metabolism;
GO: GO:0008152; P:metabolism; IEA.
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Neoptera; Endopter
NCBI_TaxID=180454;
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Bacteria; Proteobacteria; Gammaproteobacteria; Alteromonadales; Alteromonadaceae; Microbulbifer.
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                                                                      SSSGSSSSGSSSSG 15
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92.9%;
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                                                                                                                                             Score 58; DB Pred. No. 39; O; Mismatches
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                                                                                                                                                                                                                                                                                                  06332B0FCB5AD1F4
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Matches
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Q95VT3;
01-DEC-2001
01-DEC-2001
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                                                                                                                                       Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.
EMBL, AC116984; AA051376.1; -.
DictyBase; DDB0185990; CF50.
GO; GO:0003796; F:1ysozyme activity; IEA.
GO; GO:0003796; P:cell wall catabolism; IEA.
GO; GO:0016998; P:cell wall catabolism; IEA.
GO; GO:0009253; P:peptidoglycan catabolism; IEA.
InterPro; IPR002053; Glyco hydro 25.
ProDom; PD004620; Glyco hydro 25; 1.
SEQUENCE 300 AA; 30480 MW; FAEFBB7A6D967467 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=22092622; PubMed=12097910; DOI=10.1038/nature00847; Gloeckner G., Eichinger L., Szafranski K., Pachebat J., Dear P., Lehmann R., Baumgart C., Parra G., April J.F., Gulgo R., Kumpf K., Tunggal B., Cox E., Quail M.A., Platzer M., Rosenthal A., Noegel A.A.; "Sequence and analysis of chromosome 2 of Dictyostelium discoideum.";
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Q86JX6; PRELIMINARY;
01-JUN-2003 (TrEMBLrel. 24,
01-JUN-2003 (TrEMBLrel. 24,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                       Dictyostelium discoide
Eukaryota; Mycetozoa;
NCBI_TaxID=44689;
                                                       DictyBase;
SEQUENCE
                                                                                 factor have both unique and over Development 129:3657-3668(2002). EMBL; AF405695; AAL01036.1; -
                                                                                                                                                                                                                                                    Name=cf50;
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                                                                                                                        "The different components of
 Local Similarity hes 12; Conserv
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                                                       DDB0185090;
303 AA; 30
 Conservative
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                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
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cetozoa; Dictyosteliida; Dictyostelium
                                                       30854 MW;
              86.4%;
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Score 57;
Pred. No.
3; Mismatc
                                                                                                           overlapping
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Last sequence update)
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                                                                                                          a multisubunit cell number-counting overlapping functions.";
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Pred. No. 13;
                                                       2B2A3CC1CEE3CE8A CRC64;
 Mismatches
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RESULT
REPRESENTATION OF COLUMN STATE OF COLUM
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RA Adame M.D., Celniker S.E., Holt R.A., Evans C.A., Galle R.F.,
RA Mannatides P.G., Scherer S.E., Holt R.A., Evans C.A., Galle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Ballew R.M., Boyle C., Baxter E.G., Helt G., Nelson C.R., Gabler B.D.,
RA Ballew R.M., Basu A., Baxendale J., Bayrakataroglu L., Beaeley E.M.,
RA Beson K.Y., Bence P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Horil J.F., Agbayani A., An H.J., Andrews-Ffannkoch C., Baldwin D.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Gebablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Gebablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Goldek R., Gong F., Gorrell J.H., Gu Z., Gunn P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Mattei B., McIntosh T.C., McLedod M., Deymam C.,
Liu X., Mattei B., McIntosh T.C., McLedod M., McMary D.M., Nelson D.L.,
RA Mcland D.R., Nelson K.A., Nixon K., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Doulard J., Puri V., Reese M.G.,
RA Melson D.R., Nelson K.A., Stapleton M., Strong R., Shen H.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Spier E., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Ye J.,
RA Yel R.F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Ye J.,
RA Yel R.F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Ye J.,
RA Yel R.F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Xe J.,
Ra Yel R.F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Xe J.,
Ra Yel R.F., Zaveri J.S., Zhan M., Zhang G., Zhang L.,
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Genome
[3]
                                Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halpern A., Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A., George R.A., Hoskins R.A., Lavertty T., Muzny D.M., Nelson C.R., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Venter C., Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M., "Finishing a whole-genome shotgun: Release 3 of the Drosophila melanogaster euchromatic genome sequence.";

Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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25-OCT-2004 (TrEMBLrel.
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01-MAY-2000 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM
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01-JUN-2001
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Mus musculus 8 days embryo whole body cDNA, RIKEN full-length enriched library, clone:5730406M06 product:hypothetical Serine-rich region containing protein, full insert sequence. (Fragment).
Name=5730406M06Rik;
                                                                                                                                                                                                                                                                                                                                                                           FlyBase; FBgn0036949; CG7290.
GO; GO:0005576; C:extracellular; IEA.
GO; GO:0008061; F:chitin binding; IEA.
GO; GO:00080630; P:chitin metabolism; I
InterPro; IPR002557; Chitin bind PerA.
Pfam; PF01607; CBM 14; 5.
SMART; SM00494; ChEBD2; 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (DEC-2001) to the EMBL/GenBank/DDBJ EMBL; AE003514; AAF49058.3; -. EMBL; AY071500; AAL49222.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J., Champe M., Chavez C., Dorsett V., Dresnek D., Farfan D., Frise E., George R., Gonzalez M., Guarin H., Krommiller B., Li P., Liao G., Miranda A., Mungall C.J., Nunco J., Pacleb J., Paragas V., Park S., Patel S., Phouanenavong S., Wan K., Yu C., Lewis S.E., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIINE=22426069; PubMed=12537572; Milburn G.H., Camp Misra S., Crosby M.A., Mungall C.J., Milburn G.H., Prosmith C.D., Tupy J.L., Whitfied B.J., Bayraktaroglu L., Bettencourt B.R., Celniker S.E., de Grey A.D., Drysdale Harris N.L., Richter J., Russo S., Schroeder A.J., Shu Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rub
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MEDLINE=22426070; PubMed=12537573;

Kaminker J.S., Bergman C.M., Kronmiller B., Carlson J., S

Patel S., Frise B., Wheeler D.A., Lewis S.E., Rubin G.M.,

Ashburner M., Celniker S.E.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lewis S.E.;
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RP SEQUENCE FROM N.A.

RO STRAIN-C57BL/61; TISSUE=Whole body;

RA Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,

A Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,

A Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.,

RA Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Hori F.,

A Hanagaki T., Hara A., Hayatsu N., Hiramoto K., Hiraoka T., Hori F.,

A Hanagaki T., Hara A., Hayatsu N., Hiramoto K., Hiraoka T., Koto H.,

A Kawai J., Kojima Y., Konno H., Kouda M., Kasukawa T., Kato H.,

A Kawai J., Kojima Y., Konno H., Kouda M., Koya S., Kurihara C.,

A Kawai J., Kojima Y., Konno H., Kouda M., Koya S., Kurihara C.,

A Kawai J., Okido T., Owa C., Salto H., Salto R., Sakai K.,

A Matsuyama T., Miyazaki A., Nishi K., Nomura K., Numazaki R., Ohno M.,

A Matsuyama T., Okido T., Owa C., Salto H., Salto R., Sakai C., Sakai K.,

A Okazaki Y., Okido T., Owa C., Salto H., Salto R., Sakai C., Sakai K.,

A Okazaki Y., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,

A Sogabe Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

A Sogabe Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

A Muramatsu M., Hayashizaki Y., Yasunishi A., Yoshida K., Yoshino M.,

Bubai, Akol7507, Babajo77.2, -.

Babai, Akol7507, Babajo77.2, -.
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SEQUENCE FROM N.A.

STRAIN=C57BL/61; TISSUE=Whole body;

XX MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;

XX Shibata K., Itoh M., Alzawa K., Nagaoka S., Sasaki N., Carninci P.,

XX Shibata K., Itoh M., Alzawa K., Nagaoka S., Sasaki N., Carninci P.,

XX Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Harada A.,

XX Sumi N., Ishii Y., Nakamura S., Hazama M., Nishine T., Kashiwagi K.,

XX Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,

XX Fujiwake S., Inoue K., Togawa Y., Izawa M., Chara E., Watahiki M.,

XX Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,

XX Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,

XX Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,

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XX Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura T., Yoneda Y., Yoneda Y., Yoneda Y., Yoneda Y., Yoned
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MEDLINE=2049374; PubMed=11042159; DOI=10.1101/gr.145100;
Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Itoh M.,
Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
"Normalization and subtraction of cap-trapper-selected cDNAs to
prepare full-length cDNA libraries for rapid discovery of new genes.";
Genome Res. 10:1617-1630(2000).
                                                                                                                                                                                                         MGD; MGI:1913875;
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Nature 420:563-573(2002).
                                                                                                                                                                           Hypothetical
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STRAIN=C57BL/6J; TISSUE=Whole body;
MEDLINE=99279253; PubMed=1349636;
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STRAIN=C57BL/6J; TISSUE=Whole
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RL Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.

REMBL; AJ223616; CAA11491.1; -.

DR GO; GO:0016021; C:Integral to membrane; IEA.

DR GO; GO:0005524; F.Arp binding; IEA.

DR GO; GO:0005509; F:calcium ion binding; IEA.

DR GO; GO:0005509; F:calcium ion binding; IEA.

DR GO; GO:0016797; F:hydrolase activity; IEA.

DR GO; GO:0016797; F:hydrolase activity; IEA.

DR GO; GO:0016820; F:hydrolase activity, acting on acid anhydrid. ..; IEA.

DR GO; GO:0016812; P:calcium ion transport; IEA.

DR GO; GO:0016812; P:calcium ion transport; IEA.

DR GO; GO:0006812; P:calcium ion transport; IEA
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Matches 13
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STRAIN-CB1489 him-8;
STRAIN-CB1489 him-8;
MEDLINE-99134356; PubMed-9933625; DOI=10.1074/jbc.274.7.4254;
MEDLINE-99134356; PubMed-9933625; DOI=10.1074/jbc.274.7.4254;
Kraev A., Kraev N., Caratoli E.;
Kraev A., Kraev N., Caratoli E.;
"Identification and functional expression of the plasma membrane "Identification and functional expression of the plasma membrane and the plasma membrane of the plasma membrane of the plasma membrane statement of the plasma membrane of the plasma membrane and plasma plasma membrane of the plasma membrane of the plasma membrane statement of the plasma membrane of the 
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TIGRPAMS; TIGRO1517; ATPASS-IIB Ca; 1.
TIGRPAMS; TIGRO1494; ATPASS P-type; 3.
PROSITE; PS00154; ATPASE E1 E2; UNKNOWN 1.
SEQUENCE 1228 AA; 134552 MW; 1F790DFEEBBA6108 CRC64;
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01-UN-1998 (TrEMBLrel. 06, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Calcium ATPass.
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STRAIN-CB1489 him-8;
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                                                2005, 15:56:23
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